

SEQUENCE LISTING

```

<110>      Sternberg, Paul W.
           Barr, Maureen M.

<120>      POLYCYSTIC KIDNEY DISEASE GENE HOMOLOGS REQUIRED FOR MALE MATING
           BEHAVIOR IN NEMATODES AND ASSAYS BASED THEREON

<130>      18021 2901B

<140>      Unassigned
<141>      2000-09-05

<150>      09/479,467
<151>      2000-01-06

<150>      60/115,127
<151>      1999 01-06

<160>      15

<170>      Patent In Ver. 2.0

<210> 1
<211> 12912
<212> DNA
<213> Homo sapiens PKD-1 gene

<220>
<221> CDS
<222> (1)..(12912)

<400> 1
atg ccg ccc gcc ggg ccc gcc cgc ctg ggc ctg gcc ctg ggc ctg ggc 48
Met Pro Pro Ala Ala Pro Ala Arg Leu Ala Leu Ala Leu Gly Leu Gly
1 5 10 15

ctg tgg ctc ggg ggc ctg gcg ggg ggg ccc ggg cgc ggc tgc ggg ccc 96
Leu Trp Leu Gly Ala Leu Ala Gly Gly Pro Gly Arg Gly Cys Gly Pro
20 25 30

tgc gag ccc ccc tgc ctc tgc ggg cca ggc ccc ggc gcc gcc tgc cgc 144
Cys Glu Pro Pro Cys Leu Cys Gly Pro Ala Pro Gly Ala Ala Cys Arg
35 40 45

gtc aac tgc tgc ggc cgc ggg ctg cgg acg ctc ggt ccc ggc ctg cgc 192
Val Asn Cys Ser Gly Arg Gly Leu Arg Thr Leu Gly Pro Ala Leu Arg
50 55 60

atc ccc gcg gac gcc aca gag cta gac gtc tcc cac aac ctg ctc cgg 240
Ile Pro Ala Asp Ala Thr Glu Leu Asp Val Ser His Asn Leu Leu Arg
65 70 75 80

gcg ctg gac gtt ggg ctc ctg gcg aac ctc tgc gcg ctg gca gag ctg 288
Ala Leu Asp Val Gly Leu Leu Ala Asn Leu Ser Ala Leu Ala Glu Leu
85 90 95

gat ata agc aac aac aag att tct acg tta gaa gaa gga ata ttt gct 336
Asp Ile Ser Asn Asn Lys Ile Ser Thr Leu Glu Glu Gly Ile Phe Ala
100 105 110

aat tta ttt aat tta agt gaa ata aac ctg agt ggg aac ccg ttt gag 384
Asn Leu Phe Asn Leu Ser Glu Ile Asn Leu Ser Gly Asn Pro Phe Glu
115 120 125

tgt gac tgt ggc ctg gcg tgg ctg ccg caa tgg gcg gag gag cag cag 432
Cys Asp Cys Gly Leu Ala Trp Leu Pro Gln Trp Ala Glu Glu Gln Gln

```

130	gtg egg gtg gtg bag ccc gag gca gcc acg tgt gct ggg cct ggc tcc	480
Val Arg Val Val Gln Pro Glu Ala Ala Thr Cys Ala Gly Pro Gly Ser		
145	150	155
160		
ctg gct ggc bag cct ctg ctt ggc atc ccc ttg ctg gac agt ggc tgt	528	
Leu Ala Gly Gln Pro Leu Leu Gly Ile Pro Leu Leu Asp Ser Gly Cys		
165	170	175
ggc gag gag tat gtc gcc tgc ctc cct gac aac agc tca ggc acc gtg	576	
Gly Glu Glu Tyr Val Ala Cys Leu Pro Asp Asn Ser Ser Gly Thr Val		
180	185	190
200		
gca gca gtg tcc ttt tca gcc ggc cac gaa ggc ctg ctt cag cca gag	624	
Ala Ala Val Ser Phe Ser Ala Ala His Glu Gly Leu Gln Pro Glu		
195	200	205
220		
gcc tgc agc gcc ttc tgc ttc tcc acc ggc bag ggc ctc gca gcc ctc	672	
Ala Cys Ser Ala Phe Cys Phe Ser Thr Gly Gln Gly Leu Ala Ala Leu		
210	215	220
tcg gag cag ggc tgg tgc ctg tgt ggg ggc gcc bag ccc tcc agt gcc	720	
Ser Glu Gln Gly Trp Cys Leu Cys Gly Ala Ala Gln Pro Ser Ser Ala		
225	230	235
240		
tcc ttt gcc tgc ctg tcc ctc tgc tcc ggg ccc ccc gca cct cct gcc	768	
Ser Phe Ala Cys Leu Ser Leu Cys Ser Gly Pro Pro Ala Pro Pro Ala		
245	250	255
ccc acc tgt agc ggc ccc acc ctc ctc cag cac gtc ttc cct gcc tcc	816	
Pro Thr Cys Arg Gly Pro Thr Leu Leu Gln His Val Phe Pro Ala Ser		
260	265	270
cca ggg gcc acc ctg gtg ggg ccc cac gga cct ctg gcc tct ggc cag	864	
Pro Gly Ala Thr Leu Val Gly Pro His Gly Pro Leu Ala Ser Gly Gln		
275	280	285
cta gca gcc ttc cac atc gct gcc ccc ctc cct gtc act gac aca cgc	912	
Leu Ala Ala Phe His Ile Ala Ala Pro Leu Pro Val Thr Asp Thr Arg		
290	295	300
tgg gac ttc gga gac ggc tcc gcc gag gtg gat gcc gct ggg ccg gct	960	
Trp Asp Phe Gly Asp Gly Ser Ala Glu Val Asp Ala Ala Gly Pro Ala		
305	310	315
320		
gcc tcg cat cgc tat gtg ctg cct ggg cgc tat cac gtg acg gcc gtg	1008	
Ala Ser His Arg Tyr Val Leu Pro Gly Arg Tyr His Val Thr Ala Val		
325	330	335
ctg gcc ctg ggg gcc ggc tca gcc ctg ctg ggg aca gac gtg cag gtg	1056	
Leu Ala Leu Gly Ala Gly Ser Ala Leu Leu Gly Thr Asp Val Gln Val		
340	345	350
gaa gcg gca cct gcc gcc ctg gag ctc gtg tgc ccg tcc tcg gtg cag	1104	
Glu Ala Ala Pro Ala Ala Leu Glu Leu Val Cys Pro Ser Val Gln		
355	360	365
agt gac gag agc ctc gac ctc agc atc cag aac ccg ggt ggt tca ggc	1152	
Ser Asp Glu Ser Leu Asp Leu Ser Ile Gln Asn Arg Gly Gly Ser Gly		
370	375	380
ctg gag gcc gcc tac agc atc gtg gcc ctg ggc gag gag ccg gcc cga	1200	
Leu Glu Ala Ala Tyr Ser Ile Val Ala Leu Gly Glu Glu Pro Ala Arg		
385	390	395
400		
gcg gtg cac ccg ctc tgc ccc tcg gac acg gag atc ttc cct ggc aac	1248	

Ala Val His Pro	Leu Cys Pro Ser Asp Thr Glu Ile Phe Pro Gly Asn	
	405 410 415	
ggg cac tgc tac cgc ctg gtg gtg gag aag gcg gcc tgg ctg cag gcg	1296	
Gly His Cys Tyr Arg Leu Val Val Gln Lys Ala Ala Trp Leu Gln Ala		
	420 425 430	
cag gag cag tgt cag gcc tgg gcc ggg gcc gcc ctg gca atg gtg gac	1344	
Gln Glu Gln Cys Gln Ala Trp Ala Gly Ala Ala Leu Ala Met Val Asp		
	435 440 445	
agt ccc gcc gtg cag cgc ttc ctg gtc tcc cgg gtc acc agg agc cta	1392	
Ser Pro Ala Val Gln Arg Phe Leu Val Ser Arg Val Thr Arg Ser Leu		
	450 455 460	
gac gtg tgg atc ggc ttc tgg act gtg cag ggg gtg gag gtg ggc cca	1440	
Asp Val Trp Ile Gly Phe Ser Thr Val Gln Gly Val Glu Val Gly Pro		
	465 470 475 480	
ggc ccg cag ggc gag gcc ttc agc ctg gag agc tgc cag aac tgg ctg	1488	
Ala Pro Gln Gly Glu Ala Phe Ser Leu Glu Ser Cys Gln Asn Trp Leu		
	485 490 495	
ccc ggg gag cca cac cca gcc aca gcc gag cag tgc gtc cgg ctg ggg	1536	
Pro Gly Glu Pro His Pro Ala Thr Ala Glu His Cys Val Arg Leu Gly		
	500 505 510	
ccc acc ggg tgg tgt aac acc gac ctg tgc tca gcg ccg cac agc tac	1584	
Pro Thr Gly Trp Cys Asn Thr Asp Leu Cys Ser Ala Pro His Ser Tyr		
	515 520 525	
gtc tgc gag ctg cag ccc gga gcc cca gtg cag gat gcc gag aac ctg	1632	
Val Cys Glu Leu Gln Pro Gly Gly Pro Val Gln Asp Ala Glu Asn Leu		
	530 535 540	
ctc gtg gga ggc ccc agt ggg gac ctg cag gga ccc ctg acg cct ctg	1680	
Leu Val Gly Ala Pro Ser Gly Asp Leu Gln Gly Pro Leu Thr Pro Leu		
	545 550 555 560	
gca cag cag gac ggc ctc tca gcc ccg cac gag ccc gtg gag gtc atg	1728	
Ala Gln Gln Asp Gly Leu Ser Ala Pro His Glu Pro Val Glu Val Met		
	565 570 575	
gta ttc ccg ggc ctg cgt ctg agc cgt gaa gcc ttc ctc acc acg gcc	1776	
Val Phe Pro Gly Leu Arg Leu Ser Arg Glu Ala Phe Leu Thr Thr Ala		
	580 585 590	
gaa ttt ggg acc cag gag ctc cgg cgg ccc gcc cag ctg cgg ctg cag	1824	
Glu Phe Gly Thr Gln Glu Leu Arg Arg Pro Ala Gln Leu Arg Leu Gln		
	595 600 605	
gtg tac cgg ctg ctc agc aca gca ggg acc ccg gag aac ggc agc gag	1872	
Val Tyr Arg Leu Leu Ser Thr Ala Gly Thr Pro Glu Asn Gly Ser Glu		
	610 615 620	
cct gag agc agg tcc ccg gac aac agg acc cag ctg gcc ccc gcg tgc	1920	
Pro Glu Ser Arg Ser Pro Asp Asn Arg Thr Gln Leu Ala Pro Ala Cys		
	625 630 635 640	
atg cca ggg gga cgc tgg tgc cct gga gcc aac atc tgc ttg ccg ctg	1968	
Met Pro Gly Gly Arg Trp Cys Pro Gly Ala Asn Ile Cys Leu Pro Leu		
	645 650 655	
gac gcc tcc tgc cac ccc cag gcc tgc gcc aat ggc tgc acg tca ggg	2016	
Asp Ala Ser Cys His Pro Gln Ala Cys Ala Asn Gly Cys Thr Ser Gly		
	660 665 670	

cca ggg cta ccc ggg gcc ccc tat gcg cta tgg aga gag ttc ctc ttc	2064
Pro Gly Leu Pro Gly Ala Pro Tyr Ala Leu Trp Arg Glu Phe Leu Phe	
675 680 685	
ccc gtt ccc ggg ggg ccc ccc ggg cag tac tgg gtc acc ctc cag ggc	2112
Ser Val Pro Ala Gly Pro Pro Ala Gln Tyr Ser Val Thr Leu His Gly	
690 695 700	
cag gat gtc ctc atg ctc cct ggt gac ctc gtt ggc ttg cag cag gac	2160
Gln Asp Val Leu Met Leu Pro Gly Asp Leu Val Gly Leu Gln His Asp	
705 710 715 720	
gct ggc cct ggc gcc ctc ctg cag tgc tgg cgg gct ccc ggc cag cct	2208
Ala Gly Pro Gly Ala Leu Leu His Cys Ser Pro Ala Pro Gly His Pro	
725 730 735	
ggc ccc cgg gcc cgg tac ctc tcc gcc aac gcc tgg tca tgg ctg ccc	2256
Gly Pro Arg Ala Pro Tyr Leu Ser Ala Asn Ala Ser Ser Trp Leu Pro	
740 745 750	
cac ttg cca gcc cag ctg gag ggt act tgg ggc tgc cct gcc tgt gcc	2304
His Leu Pro Ala Gln Leu Glu Gly Thr Trp Gly Cys Pro Ala Cys Ala	
755 760 765	
ctg cgg ctg ctc gca caa cgg gaa cag ctc acc gtg ctg ctg ggc ttg	2352
Leu Arg Leu Leu Ala Gln Arg Glu Gln Leu Thr Val Leu Leu Gly Leu	
770 775 780	
agg ccc aac cct gga ctg cgg ctg cct ggg cgc tat gag gtc cgg gca	2400
Arg Pro Asn Pro Gly Leu Arg Leu Pro Gly Arg Tyr Glu Val Arg Ala	
785 790 795 800	
gag gtg ggc aat ggc gtg tcc agg cac aac ctc tcc tgc agc ttt gac	2448
Glu Val Gly Asn Gly Val Ser Arg His Asn Leu Ser Cys Ser Phe Asp	
805 810 815	
gtg gtc tcc cca gtg gct ggg ctg cgg gtc atc tac cct gcc ccc cgc	2496
Val Val Ser Pro Val Ala Gly Leu Arg Val Ile Tyr Pro Ala Pro Arg	
820 825 830	
gac ggc cgc ctc tac gtg ccc acc aac ggc tca gcc ttg gtg ctc cag	2544
Asp Gly Arg Leu Tyr Val Pro Thr Asn Gly Ser Ala Leu Val Leu Gln	
835 840 845	
gtg gac tct ggt gcc aac gcc acg gcc acg gct cgc tgg cct ggg ggc	2592
Val Asp Ser Gly Ala Asn Ala Thr Ala Thr Ala Arg Trp Pro Gly Gly	
850 855 860	
agt ctc agc gcc cgc ttt gag aat gtc tgc cct gcc ctg gtg gcc acc	2640
Ser Leu Ser Ala Arg Phe Glu Asn Val Cys Pro Ala Leu Val Ala Thr	
865 870 875 880	
ttc gtg ccc gcc tgc ccc tgg gag acc aac gat acc ctg ttc tca gtg	2688
Phe Val Pro Ala Cys Pro Trp Glu Thr Asn Asp Thr Leu Phe Ser Val	
885 890 895	
gta gca ctg ccg tgg ctc agt gag ggg gag cac gtg gtg gac gtg gtg	2736
Val Ala Leu Pro Trp Leu Ser Glu Gly Glu His Val Val Asp Val Val	
900 905 910	
gtg gaa aac agc gcc agc cgg gcc aac ctc agc ctg cgg gtg acg gcg	2784
Val Glu Asn Ser Ala Ser Arg Ala Asn Leu Ser Leu Arg Val Thr Ala	
915 920 925	
gag gag ccc atc tgt ggc ctc cgc gcc acg ccc agc ccc gag gcc cgt	2832
Glu Glu Pro Ile Cys Gly Leu Arg Ala Thr Pro Ser Pro Glu Ala Arg	
930 935 940	

gta ctg cag gga gtc cta gtg agg tac agc ccc gtg gtg gag gcc ggc	2880
Val Leu Gln Gly Val Leu Val Arg Tyr Ser Pro Val Val Gln Ala Gly	
945 950 955 960	
tcg gac atg gtc ttc cgg tgg acc atc aac gac aag cag tcc ctg acc	2923
Ser Asp Met Val Phe Arg Trp Thr Ile Asn Asp Lys Gln Ser Leu Thr	
965 970 975	
ttc cag aac gtg gtc ttc aat gtc att tac cag agc ggc ggc gtc ttc	2976
Phe Gln Asn Val Val Phe Asn Val Ile Tyr Gln Ser Ala Val Phe	
980 985 990	
aag ctc tca ctg acg gcc tcc aac cac gtg agc aac gtc acc gtg aac	3024
Lys Leu Ser Leu Thr Ala Ser Asn His Val Ser Asn Val Thr Val Asn	
995 1000 1005	
tac aac gta acc gtg gag cgg atg aac agg atg cag ggt ctg cag gtc	3072
Tyr Asn Val Thr Val Gln Arg Met Asn Arg Met Gln Gly Leu Gln Val	
1010 1015 1020	
tcc aca gtg cgg gcc gtg ctg tcc ccc aat gcc acg cta gca ctg acc	3120
Ser Thr Val Pro Ala Val Leu Ser Pro Asn Ala Thr Leu Ala Leu Thr	
1025 1030 1035 1040	
ggc gcc gtg ctg gtg gac tcg gcc gtg gag gtg gcc ttc ctg tgg acc	3163
Ala Gly Val Leu Val Asp Ser Ala Val Glu Val Ala Phe Leu Trp Thr	
1045 1050 1055	
ttt ggg gat ggg gag cag gcc ctc cac cag ttc cag cct cgg tac aac	3216
Phe Gly Asp Gly Glu Gln Ala Leu His Gln Phe Gln Pro Pro Tyr Asn	
1060 1065 1070	
gag tcc ttc cca gtt cca gac ccc tcg gtg gcc cag gtg ctg gtg gag	3264
Glu Ser Phe Pro Val Pro Asp Pro Ser Val Ala Gln Val Leu Val Glu	
1075 1080 1085	
cac aat gtc acg cac acc tac gct gcc cca ggt gag tac ctc ctg acc	3312
His Asn Val Thr His Thr Tyr Ala Ala Pro Gly Glu Tyr Leu Leu Thr	
1090 1095 1100	
gtg ctg gca tct aat gcc ttc gag aac ctg acg cag cag gtg cct gtg	3360
Val Leu Ala Ser Asn Ala Phe Glu Asn Leu Thr Gln Gln Val Pro Val	
1105 1110 1115 1120	
agc gtg cgc gcc tcc ctg ccc tcc gtg gct gtg ggt gtg agt gac ggc	3408
Ser Val Arg Ala Ser Leu Pro Ser Val Ala Val Gly Val Ser Asp Gly	
1125 1130 1135	
gtc ctg gtg gcc ggc cgg ccc gtc acc ttc tac cgg cac cgg ctg ccc	3456
Val Leu Val Ala Gly Arg Pro Val Thr Phe Tyr Pro His Pro Leu Pro	
1140 1145 1150	
tcg cct ggg ggt gtt ctt tac acg tgg gac ttc ggg gac gcc tcc cct	3504
Ser Pro Gly Gly Val Leu Tyr Thr Trp Asp Phe Gly Asp Gly Ser Pro	
1155 1160 1165	
gtc ctg acc cag agc cag ccg gct gcc aac cac acc tat gcc tcg agg	3552
Val Leu Thr Gln Ser Gln Pro Ala Ala Asn His Thr Tyr Ala Ser Arg	
1170 1175 1180	
ggc acc tac cac gtg cgc ctg gag gtc aac aac acg gtg agc ggt ggc	3600
Gly Thr Tyr His Val Arg Leu Glu Val Asn Asn Thr Val Ser Gly Ala	
1185 1190 1195 1200	
gcg gcc cag gcg gat gtg cgc gtc ttt gag gag ctc cgc gga ctc agc	3648
Ala Ala Gln Ala Asp Val Arg Val Phe Glu Glu Leu Arg Gly Leu Ser	

1205	1210	1215	
gtg gac atg agc ctg gcc gtg gag cag ggc gcc ccc gtg gtg gtc agc Val Asp Met Ser Leu Ala Val Glu Gln Gly Ala Pro Val Val Val Ser 1220 1225 1230			3696
gcc ggc gtg cag acg ggc gac aac atc acg tgg acc ttc gac atg ggg Ala Ala Val Gln Thr Gly Asp Asn Ile Thr Trp Thr Phe Asp Met Gly 1235 1240 1245			3744
gac gcc acc gtg ctg tcc ggc ccg gag gca aca gtg gag cat gtg tac Asp Gly Thr Val Leu Ser Gly Pro Glu Ala Thr Val Glu His Val Tyr 1250 1255 1260			3792
ctg cgg gca cag aac tgc aca gtg acc gtg ggt ggc gcc agc ccc gcc Leu Arg Ala Gln Asn Cys Thr Val Thr Val Gly Ala Gly Ser Pro Ala 1265 1270 1275 1280			3840
ggc cac ctg gcc cgg agc ctg cac gtg ctg gtc ttc gtc ctg gag gtg Gly His Leu Ala Arg Ser Leu His Val Leu Val Phe Val Leu Glu Val 1285 1290 1295			3888
ctg cgt gtt gaa ccc gcc gcc tgc atc ccc acg cag cct gac ggc cgg Leu Arg Val Glu Pro Ala Ala Cys Ile Pro Thr Gln Pro Asp Ala Arg 1300 1305 1310			3936
ctc acg gcc tac gtc acc ggg aac ccg gcc cac tac ctc ttc gac tgg Leu Thr Ala Tyr Val Thr Gly Asn Pro Ala His Tyr Leu Phe Asp Trp 1315 1320 1325			3984
acc ttc ggg gat ggc tcc tcc aac acg acc gtg cgg ggg tgc ccg acg Thr Phe Gly Asp Gly Ser Ser Asn Thr Thr Val Arg Gly Cys Pro Thr 1330 1335 1340			4032
gtg aca cac aac ttc acg cgg agc ggc acg ttc ccc ctg ggc ctg gtg Val Thr His Asn Phe Thr Arg Ser Gly Thr Phe Pro Leu Ala Leu Val 1345 1350 1355 1360			4080
ctg tcc agc cgc gtg aac agg gcg cat tac ttc acc agc atc tgc gtg Leu Ser Ser Arg Val Asn Arg Ala His Tyr Phe Thr Ser Ile Cys Val 1365 1370 1375			4128
gag cca gag gtg ggc aac gtc acc ctg cag cca gag agg cag ttt gtg Glu Pro Glu Val Gly Asn Val Thr Leu Gln Pro Glu Arg Gln Phe Val 1380 1385 1390			4176
cag ctc ggg gac gag gcc tgg ctg gtg gca tgt gcc tgg ccc ccg ttc Gln Leu Gly Asp Glu Ala Trp Leu Val Ala Cys Ala Trp Pro Pro Phe 1395 1400 1405			4224
ccc tac cgc tac acc tgg gac ttt ggc acc gag gaa gcc gcc ccc acc Pro Tyr Arg Tyr Thr Trp Asp Phe Gly Thr Glu Glu Ala Ala Pro Thr 1410 1415 1420			4272
cgt gcc agg ggc cct gag gtg acg ttc atc tac cga gac cca ggc tcc Arg Ala Arg Gly Pro Glu Val Thr Phe Ile Tyr Arg Asp Pro Gly Ser 1425 1430 1435 1440			4320
tat ctt gtg aca gtc acc gcg tcc aac aac atc tct gct gcc aat gac Tyr Leu Val Thr Val Thr Ala Ser Asn Asn Ile Ser Ala Ala Asn Asp 1445 1450 1455			4368
tca gcc ctg gtg gag gtg cag gag ccc gtg ctg gtc acc agc atc aag Ser Ala Leu Val Glu Val Gln Glu Pro Val Leu Val Thr Ser Ile Lys 1460 1465 1470			4416
gtc aat ggc tcc ctt ggg ctg gag ctg cag cag ccg tac ctg ttc tct			4464

Val Asn Gly Ser Leu Gly Leu Glu Leu Gln Gln Pro Tyr Leu Phe Ser	
1475 1480 1435	
ggt gtg ggc cgt ggg cgc ccc gcc agc tac ctg tgg gat ctg ggg gac	4512
Ala Val Gly Arg Gly Arg Pro Ala Ser Tyr Leu Trp Asp Leu Gly Asp	
1490 1495 1500	
ggt ggg tgg ctc gag ggt ccg gag gtc acc cac gct tac aac agc aca	4550
Gly Gly Trp Leu Glu Gly Pro Glu Val Thr His Ala Tyr Asn Ser Thr	
1505 1510 1515 1520	
ggt gac ttc acc gtt agg gtg gcc ggc tgg aat gag gtg agc cgc agc	4608
Gly Asp Phe Thr Val Arg Val Ala Gly Trp Asn Glu Val Ser Arg Ser	
1525 1530 1535	
gag gcc tgg ctc aat gtg acg gtg aag cgg cgc gtg cgg ggg ctc gtc	4656
Glu Ala Trp Leu Asn Val Thr Val Lys Arg Arg Val Arg Gly Leu Val	
1540 1545 1550	
gtc aat gca agc cgc acg gtg gtg ccc ctg aat ggg agc gtg agc ttc	4704
Val Asn Ala Ser Arg Thr Val Val Pro Leu Asn Gly Ser Val Ser Phe	
1555 1560 1565	
agc acg tgg ctg gag gcc ggc agt gat gtg cgc tat tcc tgg gtg ctc	4752
Ser Thr Ser Leu Glu Ala Gly Ser Asp Val Arg Tyr Ser Trp Val Leu	
1570 1575 1580	
tgt gac cgc tgc acg ccc atc cct ggg ggt cct acc atc tct tac acc	4800
Cys Asp Arg Cys Thr Pro Ile Pro Gly Gly Pro Thr Ile Ser Tyr Thr	
1585 1590 1595 1600	
ttc cgc tcc gtg ggc acc ttc aat atc atc gtc acg gct gag aac gag	4848
Phe Arg Ser Val Gly Thr Phe Asn Ile Ile Val Thr Ala Glu Asn Glu	
1605 1610 1615	
gtg ggc tcc gcc cag gac agc atc ttc gtc tat gtc ctg cag ctc ata	4896
Val Gly Ser Ala Gln Asp Ser Ile Phe Val Tyr Val Leu Gln Leu Ile	
1620 1625 1630	
gag ggg ctg cag gtg gtg ggc ggt ggc cgc tac ttc ccc acc aac cac	4944
Glu Gly Leu Gln Val Val Gly Gly Gly Arg Tyr Phe Pro Thr Asn His	
1635 1640 1645	
acg gta cag ctg cag gcc gtg gtt agg gat ggc acc aac gtc tcc tac	4992
Thr Val Gln Leu Gln Ala Val Val Arg Asp Gly Thr Asn Val Ser Tyr	
1650 1655 1660	
agc tgg act gcc tgg agg gac agg ggc ccg gcc ctg gcc ggc agc ggc	5040
Ser Trp Thr Ala Trp Arg Asp Arg Gly Pro Ala Leu Ala Gly Ser Gly	
1665 1670 1675 1680	
aaa ggc ttc tgg ctc acc gtg ctc gag gcc ggc acc tac cat gtg cag	5088
Lys Gly Phe Ser Leu Thr Val Leu Glu Ala Gly Thr Tyr His Val Gln	
1685 1690 1695	
ctg cgg gcc acc aac atg ctg ggc agc gcc tgg gcc gac tgc acc atg	5136
Leu Arg Ala Thr Asn Met Leu Gly Ser Ala Trp Ala Asp Cys Thr Met	
1700 1705 1710	
gac ttc gtg gag cct gtg ggg tgg ctg atg gtg gcc gcc tcc ccg aac	5184
Asp Phe Val Glu Pro Val Gly Trp Leu Met Val Ala Ala Ser Pro Asn	
1715 1720 1725	
cca gct gcc gtc aac aca agc gtc acc ctc agt gcc gag ctg gct ggt	5232
Pro Ala Ala Val Asn Thr Ser Val Thr Leu Ser Ala Glu Leu Ala Gly	
1730 1735 1740	

ggc agt ggt gtc gta tac act tgg tcc ttg gag gag ggg ctg agc tgg Gly Ser Gly Val Val Tyr Thr Trp Ser Leu Glu Glu Gly Leu Ser Trp 1745 1750 1755 1760	5280
gag acc tcc gag cca ttt acc acc cat agt ttc ccc aca ccc ggc ctg Glu Thr Ser Glu Pro Phe Thr Thr His Ser Phe Pro Thr Pro Gly Leu 1765 1770 1775	5328
cac ttg gtc acc atg aag gca ggg aac cag ctg ggc tca gcc aac gcc His Leu Val Thr Met Thr Ala Gly Asn Pro Leu Gly Ser Ala Asn Ala 1780 1785 1790	5376
acc gtg gaa gtg gat gtg cag gtg cct gtg agt ggc ctg agc atc agg Thr Val Glu Val Asp Val Gln Val Pro Val Ser Gly Leu Ser Ile Arg 1795 1800 1805	5424
gcc agc gag ccc gga ggc agc ttc gtg ggc gcc ggg tcc tct gtg ccc Ala Ser Glu Pro Gly Gly Ser Phe Val Ala Ala Gly Ser Ser Val Pro 1810 1815 1820	5472
ttt tgg ggg cag ctg gcc aag ggc acc aat gtg agc tgg tgc tgg gct Phe Trp Gly Gln Leu Ala Thr Gly Thr Asn Val Ser Trp Cys Trp Ala 1825 1830 1835 1840	5520
gtg ccc ggc ggt agc agc aag cgt ggc cct cat gtc acc atg gtc ttc Val Pro Gly Gly Ser Ser Lys Arg Gly Pro His Val Thr Met Val Phe 1845 1850 1855	5568
cag gat gct ggc acc ttc tcc atc cgg ctg aat gcc tcc aac gca gtc Pro Asp Ala Gly Thr Phe Ser Ile Arg Leu Asn Ala Ser Asn Ala Val 1860 1865 1870	5616
agc tgg gtc tca gcc acg tac aac ctg aag gag gag gag ccc atc gtg Ser Trp Val Ser Ala Thr Tyr Asn Leu Thr Ala Glu Glu Pro Ile Val 1875 1880 1885	5664
ggc ctg gtg ctg tgg gcc agc agc aag gtg gtg ggc ccc ggg cag ctg Gly Leu Val Leu Trp Ala Ser Ser Lys Val Val Ala Pro Gly Gln Leu 1890 1895 1900	5712
gtc cat ttt cag atc ctg ctg gct gcc ggc tca gct gtc acc ttc cgc Val His Phe Gln Ile Leu Leu Ala Ala Gly Ser Ala Val Thr Phe Arg 1905 1910 1915 1920	5760
cta cag gtc ggc ggg gcc aac ccc gag gtg ctg ccc ggg ccc cgt ttc Leu Gln Val Gly Gly Ala Asn Pro Glu Val Leu Pro Gly Pro Arg Phe 1925 1930 1935	5808
tcc cac agc ttc ccc cgc gtc gga gac cac gtg gtg agc gtg cgg ggc Ser His Ser Phe Pro Arg Val Gly Asp His Val Val Ser Val Arg Gly 1940 1945 1950	5856
aaa aac cac gtg agc tgg gcc cag ggc cag gtg cgc atc gtg gtg ctg Lys Asn His Val Ser Trp Ala Gln Ala Gln Val Arg Ile Val Val Leu 1955 1960 1965	5904
gag gcc gtg agt ggg ctg cag gtg ccc aac tgc tgc gag cct ggc atc Glu Ala Val Ser Gly Leu Gln Val Pro Asn Cys Cys Glu Pro Gly Ile 1970 1975 1980	5952
gcc acg ggc act gag agg aac ttc aca gcc cgc gtg cag cgc ggc tct Ala Thr Gly Thr Glu Arg Asn Phe Thr Ala Arg Val Gln Arg Gly Ser 1985 1990 1995 2000	6000
cgg gtc gcc tac gcc tgg tac ttc tgc ctg cag aag gtc cag ggc gac Arg Val Ala Tyr Ala Trp Tyr Phe Ser Leu Gln Lys Val Gln Gly Asp 2005 2010 2015	6048

tgg ctg gtc atc ctg tgg ggc cgc gac gtc acc tac acg ccc gtg gcc Ser Leu Val Ile Leu Ser Gly Arg Asp Val Thr Tyr Thr Pro Val Ala 2020 2025 2030	5096
ggg ggg ctg ttg gag atc cag gtg cgc gcc ttc aac gcc ctg ggc agt Ala Gly Leu Leu Glu Ile Gln Val Arg Ala Phe Asn Ala Leu Gly Ser 2035 2040 2045	5144
gag aac cgc acg ctg gtg ctg gag gtc cag gac gcc gtc cag tat gtg Glu Asn Arg Thr Leu Val Leu Glu Val Gln Asp Ala Val Gln Tyr Val 2050 2055 2060	5192
gcc ctg cag agc ggc ccc cgc ttc acc aac cgc tgg ggc cag ttt gag Ala Leu Gln Ser Gly Pro Cys Phe Thr Asn Arg Ser Ala Gln Phe Glu 2065 2070 2075 2080	5240
gcc gcc acc agc ccc agc ccc cgg cgt gtg gcc tac cag tgg gac ttt Ala Ala Thr Ser Pro Ser Pro Arg Arg Val Ala Tyr His Trp Asp Phe 2085 2090 2095	5288
ggg gat ggg tgg cca ggg cag gac aca gat gag ccc agg gcc gag cac Gly Asp Gly Ser Pro Gly Gln Asp Thr Asp Glu Pro Arg Ala Glu His 2100 2105 2110	5336
ccc tac ctg agg cct ggg gac tac cgc gtg cag gtg aac gcc tcc aac Ser Tyr Leu Arg Pro Gly Asp Tyr Arg Val Gln Val Asn Ala Ser Asn 2115 2120 2125	5384
ctg gtg agc ttc ttc gtg ggc cag gcc acg gtg acc gtc cag gtg ctg Leu Val Ser Phe Phe Val Ala Gln Ala Thr Val Thr Val Gln Val Leu 2130 2135 2140	5432
gcc tgc cgg gag cgg gag gtg gac gtg gtc ctg ccc ctg cag gtg ctg Ala Cys Arg Glu Pro Glu Val Asp Val Val Leu Pro Leu Gln Val Leu 2145 2150 2155 2160	5480
atg cgg cga tca cag cgc aac tac ttg gag gcc cac gtt gac ctg cgc Met Arg Arg Ser Gln Arg Asn Tyr Leu Glu Ala His Val Asp Leu Arg 2165 2170 2175	5528
gac tgc gtc acc tac cag act gag tac cgc tgg gag gtg tat cgc acc Asp Cys Val Thr Tyr Gln Thr Glu Tyr Arg Trp Glu Val Tyr Arg Thr 2180 2185 2190	5576
gcc agc tgc cag cgg cgg ggg cgc cca ggc cgt gtg gcc ctg ccc ggc Ala Ser Cys Gln Arg Pro Gly Arg Pro Ala Arg Val Ala Leu Pro Gly 2195 2200 2205	5624
gtg gac gtg agc cgg cct cgg ctg gtg ctg cgg cgg ctg ggc ctg cct Val Asp Val Ser Arg Pro Arg Leu Val Leu Pro Arg Leu Ala Leu Pro 2210 2215 2220	5672
gtg ggg cac tac tgc ttt gtg ttt gtc gtg tca ttt ggg gac acg cca Val Gly His Tyr Cys Phe Val Phe Val Val Ser Phe Gly Asp Thr Pro 2225 2230 2235 2240	5720
ctg aca cag agc atc cag gcc aat gtg acg gtg gcc ccc gag cgc ctg Leu Thr Gln Ser Ile Gln Ala Asn Val Thr Val Ala Pro Glu Arg Leu 2245 2250 2255	5768
gtg ccc atc att gag ggt ggc tca tac cgc gtg tgg tca gac aca cgg Val Pro Ile Ile Glu Gly Gly Ser Tyr Arg Val Trp Ser Asp Thr Arg 2260 2265 2270	5816
gac ctg gtg ctg gat ggg agc gag tcc tac gac ccc aac ctg gag gac Asp Leu Val Leu Asp Gly Ser Glu Ser Tyr Asp Pro Asn Leu Glu Asp 2275 2280 2285 2290	5864

2275	2280	2285	
ggc gac cag acg ccg ctc agt ttc cac tgg gcc tgt gtg gat tag aca			6912
Gly Asp Gln Thr Pro Leu Ser Phe His Trp Ala Cys Val Ala Ser Thr			
2290	2295	2300	
cag agg gag gat ggc ggg tgt ggg ctg aac ttc ggg ccc cgc ggg agc			6950
Gln Arg Glu Ala Gly Gly Cys Ala Leu Asn Phe Gly Pro Arg Gly Ser			
2305	2310	2315	2320
agg acg gtc acc att cca cgg gag cgg ctg ggg gat ggc gtg gag tac			7008
Ser Thr Val Thr Ile Pro Arg Glu Arg Leu Ala Ala Gly Val Glu Tyr			
2325	2330	2335	
acc ttc agc ctg acc gtg tgg aag gcc ggc cgc aag gag gag gcc acc			7056
Thr Phe Ser Leu Thr Val Trp Lys Ala Gly Arg Lys Glu Glu Ala Thr			
2340	2345	2350	
aac cag acg gtc ctg att cgg agt ggc cgg gtc ccc att gtc tcc ttg			7104
Asn Gln Thr Val Leu Ile Arg Ser Gly Arg Val Pro Ile Val Ser Leu			
2355	2360	2365	
gag tgt gtc tcc tgc aag gca cag gcc gtc tac gaa gtc agc cgc agc			7152
Glu Cys Val Ser Cys Lys Ala Gln Ala Val Tyr Glu Val Ser Arg Ser			
2370	2375	2380	
tcc tac gtc tac ttg gag gcc cgc tgc ctc aat tgc agc agc gcc tcc			7200
Ser Tyr Val Tyr Leu Glu Gly Arg Cys Leu Asn Cys Ser Ser Gly Ser			
2385	2390	2395	2400
aag cga ggg cgg tgg gct gca cgt acc ttc agc aac aag acc ctg gtc			7248
Lys Arg Gly Arg Trp Ala Ala Arg Thr Phe Ser Asn Lys Thr Leu Val			
2405	2410	2415	
ctg gat gag acc acc aca tcc acg gcc agt gca gcc atg cga ctg gtc			7296
Leu Asp Glu Thr Thr Thr Ser Thr Gly Ser Ala Gly Met Arg Leu Val			
2420	2425	2430	
ctg cgg cgg gcc gtc ctg cgg gac gcc gag gga tac acc ttc acc ctc			7344
Leu Arg Arg Gly Val Leu Arg Asp Gly Glu Gly Tyr Thr Phe Thr Leu			
2435	2440	2445	
acg gtc ctg gcc cgc tct gcc gag gag gag gcc tgc gcc tcc atc cgc			7392
Thr Val Leu Gly Arg Ser Gly Glu Glu Glu Gly Cys Ala Ser Ile Arg			
2450	2455	2460	
ctg tcc ccc aac cgc ccg ccg ctg ggg gcc tct tgc cgc ctc ttc cca			7440
Leu Ser Pro Asn Arg Pro Pro Leu Gly Gly Ser Cys Arg Leu Phe Pro			
2465	2470	2475	2480
ctg gcc gct gtc cac gcc ctc acc acc aag gtc cac ttc gaa tgc acc			7488
Leu Gly Ala Val His Ala Leu Thr Thr Lys Val His Phe Glu Cys Thr			
2485	2490	2495	
ggc tgg cat gac gcg gag gat gct gcc gcc ccg ctg gtc tac gcc ctg			7536
Gly Trp His Asp Ala Glu Asp Ala Gly Ala Pro Leu Val Tyr Ala Leu			
2500	2505	2510	
ctg ctg cgg cgc tgt cgc cag gcc cac tgc gag gag ttc tgt gtc tac			7584
Leu Leu Arg Arg Cys Arg Gln Gly His Cys Glu Glu Phe Cys Val Tyr			
2515	2520	2525	
aag gcc agc ctc tcc agc tac gga gcc gtc ctg ccc ccg ggt ttc agg			7632
Lys Gly Ser Leu Ser Ser Tyr Gly Ala Val Leu Pro Pro Gly Phe Arg			
2530	2535	2540	
cca cac ttc gag gtc gcc ctg gcc gtc gtc gtc cag gac cag ctg gga			7680

Pro His Phe Glu Val Gly Leu Ala Val Val Val Gln Asp Gln Leu Gly	
2545 2550 2555 2560	
gcc ggt gtg gtc gcc ctc aac agg tct ttg gcc atc acc ctc cca gag	7723
Ala Ala Val Val Ala Leu Asn Arg Ser Leu Ala Ile Thr Leu Pro Glu	
2565 2570 2575	
ccc aac ggc agc gca acg ggg ctc aca gtc tgg ctg cac ggg ctc acc	7776
Pro Asn Gly Ser Ala Thr Gly Leu Thr Val Trp Leu His Gly Leu Thr	
2580 2585 2590	
gct agt gtg ctc cca ggg ctg ctg cgg cag gcc gat ccc cag cac gtc	7824
Ala Ser Val Leu Pro Gly Leu Leu Arg Gln Ala Asp Pro Gln His Val	
2595 2600 2605	
atc gag tac tgc ttg gcc ctg gtc acc gtg ctg aac gag tac gag cgg	7872
Ile Glu Tyr Ser Leu Ala Leu Val Thr Val Leu Asn Glu Tyr Glu Arg	
2610 2615 2620	
gcc ctg gac gtg gcg gca gag ccc aag cac gag cgg cag cac cga gcc	7920
Ala Leu Asp Val Ala Ala Glu Pro Lys His Glu Arg Gln His Arg Ala	
2625 2630 2635 2640	
cag ata cgc aag aac atc acg gag act ctg gtg tcc ctg agg gtc cac	7968
Gln Ile Arg Lys Asn Ile Thr Glu Thr Leu Val Ser Leu Arg Val His	
2645 2650 2655	
act gtg gat gac atc cag cag atc gct gct gcg ctg gcc cag tgc atg	8016
Thr Val Asp Asp Ile Gln Gln Ile Ala Ala Ala Leu Ala Gln Cys Met	
2660 2665 2670	
ggg ccc agc agg gag ctc gta tgc cgc tgc tgc ctg aag cag acg ctg	8064
Gly Pro Ser Arg Glu Leu Val Cys Arg Ser Cys Leu Lys Gln Thr Leu	
2675 2680 2685	
cac aag ctg gag gcc atg atg ctc atc ctg cag gca gag acc acc gcg	8112
His Lys Leu Glu Ala Met Met Leu Ile Leu Gln Ala Glu Thr Thr Ala	
2690 2695 2700	
ggc acc gtg acg ccc acc gcc atc gga gac agc atc ctc aac atc aca	8160
Gly Thr Val Thr Pro Thr Ala Ile Gly Asp Ser Ile Leu Asn Ile Thr	
2705 2710 2715 2720	
gga gac ctc atc cac ctg gcc agc tgc gac gtg cgg gca cca cag ccc	8208
Gly Asp Leu Ile His Leu Ala Ser Ser Asp Val Arg Ala Pro Gln Pro	
2725 2730 2735	
tca gag ctg gga gcc gag tca cca tct cgg atg gtg gcg tcc cag gcc	8256
Ser Glu Leu Gly Ala Glu Ser Pro Ser Arg Met Val Ala Ser Gln Ala	
2740 2745 2750	
tac aac ctg acc tct gcc ctc atg cgc atc ctc atg cgc tcc cgc gtg	8304
Tyr Asn Leu Thr Ser Ala Leu Met Arg Ile Leu Met Arg Ser Arg Val	
2755 2760 2765	
ctc aac gag gag ccc ctg acg ctg gcg ggc gag gag atc gtg gcc cag	8352
Leu Asn Glu Glu Pro Leu Thr Leu Ala Gly Glu Glu Ile Val Ala Gln	
2770 2775 2780	
ggc aag cgc tgc gac ccg cgg agc ctg ctg tgc tat ggc ggc gcc cca	8400
Gly Lys Arg Ser Asp Pro Arg Ser Leu Leu Cys Tyr Gly Gly Ala Pro	
2785 2790 2795 2800	
ggg cct ggc tgc cac ttc tcc atc ccc gag gct ttc agc ggg gcc ctg	8448
Gly Pro Gly Cys His Phe Ser Ile Pro Glu Ala Phe Ser Gly Ala Leu	
2805 2810 2815	

gac aac ctc agt gac gtg gtg cag ctc atc ttt ctg gtg gac tcc aat Ala Asn Leu Ser Asp Val Val Gln Leu Ile Phe Leu Val Asp Ser Asn 2820 2825 2830	8496
ccc ttt ccc ttt ggc tat atc agc aac tac acc gtc tcc acc aag gtg Pro Phe Pro Phe Gly Tyr Ile Ser Asn Tyr Thr Val Ser Thr Lys Val 2835 2840 2845	8544
ggc tog atg gca ttc cag aca cag gcc ggc gcc cag atc ccc atc gag Ala Ser Met Ala Phe Gln Thr Gln Ala Gly Ala Gln Ile Pro Ile Glu 2850 2855 2860	8592
egg ctg gcc tca gag cgc gcc acc acc gtg aag gtg ccc aac aac tog Arg Leu Ala Ser Glu Arg Ala Ile Thr Val Lys Val Pro Asn Asn Ser 2865 2870 2875 2880	8640
gac tgg gct gcc egg ggc cac cgc agc tcc gcc aac tcc gcc aac tcc Asp Trp Ala Ala Arg Gly His Arg Ser Ser Ala Asn Ser Ala Asn Ser 2885 2890 2895	8688
gtt gtg gtc cag ccc cag gcc tcc gtc ggt gct gtg gtc acc ctg gac Val Val Val Gln Pro Gln Ala Ser Val Gly Ala Val Val Thr Leu Asp 2900 2905 2910	8736
agc agc aac cct ggc gcc ggg ctg cat ctg cag ctc aac tat acg ctg Ser Ser Asn Pro Ala Ala Gly Leu His Leu Gln Leu Asn Tyr Thr Leu 2915 2920 2925	8784
ctg gac ggc cac tac ctg tct gag gaa cct gag ccc tac ctg gca gtc Leu Asp Gly His Tyr Leu Ser Glu Glu Pro Glu Pro Tyr Leu Ala Val 2930 2935 2940	8832
tac cta cac tog gag ccc cgg ccc aat gag cac aac tgc tog gct agc Tyr Leu His Ser Glu Pro Arg Pro Asn Glu His Asn Cys Ser Ala Ser 2945 2950 2955 2960	8880
agg agg atc cgc cca gag tca ctc cag ggt gct gac cac cgg ccc tac Arg Arg Ile Arg Pro Glu Ser Leu Gln Gly Ala Asp His Arg Pro Tyr 2965 2970 2975	8928
acc ttc ttc att tcc cgg ggg agc aga gac cca gcg ggg agt tac cat Thr Phe Phe Ile Ser Pro Gly Ser Arg Asp Pro Ala Gly Ser Tyr His 2980 2985 2990	8976
ctg aac ctc tcc agc cac ttc cgc tgg tgc gcg ctg cag gtg tcc gtg Leu Asn Leu Ser Ser His Phe Arg Trp Ser Ala Leu Gln Val Ser Val 2995 3000 3005	9024
ggc ctg tac acg tcc ctg tgc cag tac ttc agc gag gag gac atg gtg Gly Leu Tyr Thr Ser Leu Cys Gln Tyr Phe Ser Glu Glu Asp Met Val 3010 3015 3020	9072
tgg cgg aca gag ggg ctg ctg ccc ctg gag gag acc tgc ccc cgc cag Trp Arg Thr Glu Gly Leu Leu Pro Leu Glu Glu Thr Ser Pro Arg Gln 3025 3030 3035 3040	9120
gcc gtc tgc ctc acc cgc cac ctc acc gcc ttc ggc gcc agc ctc ttc Ala Val Cys Leu Thr Arg His Leu Thr Ala Phe Gly Ala Ser Leu Phe 3045 3050 3055	9168
gtg ccc cca agc cat gtc cgc ttt gtg ttt cct gag ccg aca gcg gat Val Pro Pro Ser His Val Arg Phe Val Phe Pro Glu Pro Thr Ala Asp 3060 3065 3070	9216
gta aac tac atc gtc atg ctg aca tgt gct gtg tgc ctg gtg acc tac Val Asn Tyr Ile Val Met Leu Thr Cys Ala Val Cys Leu Val Thr Tyr 3075 3080 3085	9264

atg gtc atg gcc gcc atc ctg cac aag ctg gac cag ttg gat gcc agc Met Val Met Ala Ala Ile Leu His Lys Leu Asp Gln Leu Asp Ala Ser 3090 3095 3100	9312
egg gcc egg gcc atc cct ttc tgt ggg cag egg gcc egg ttc aag tac Arg Gly Arg Ala Ile Pro Phe Cys Gly Gln Arg Gly Arg Phe Lys Tyr 3105 3110 3115 3120	9360
gag atc ctg gtc aag aca ggc tgg ggc cgg gcc tca ggt acc acg gcc Glu Ile Leu Val Lys Thr Gly Trp Gly Arg Gly Ser Gly Thr Thr Ala 3125 3130 3135	9408
cac gtg gcc atc atg ctg tat ggg gtg gac agc egg agc gcc cac egg His Val Gly Ile Met Leu Tyr Gly Val Asp Ser Arg Ser Gly His Arg 3140 3145 3150	9456
cac ctg gac gcc gac aga gcc ttc cac cgc aac agc ctg gac atc ttc His Leu Asp Gly Asp Arg Ala Phe His Arg Asn Ser Leu Asp Ile Phe 3155 3160 3165	9504
egg atc gcc acc cgg cac agc ctg ggt agc gtg tgg aag atc cga gtg Arg Ile Ala Thr Pro His Ser Leu Gly Ser Val Trp Lys Ile Arg Val 3170 3175 3180	9552
tgg cac gac aac aaa ggg ctg agc cct gcc tgg ttc ctg cag cac gtc Trp His Asp Asn Lys Gly Leu Ser Pro Ala Trp Phe Leu Gln His Val 3185 3190 3195 3200	9600
atc gtc agg gac ctg cag acg gca cgc agc gcc ttc ttc ctg gtc aat Ile Val Arg Asp Leu Gln Thr Ala Arg Ser Ala Phe Phe Leu Val Asn 3205 3210 3215	9648
gac tgg ctt tgg gtg gag acg gag gcc aac ggg gcc ctg gtg gag aag Asp Trp Leu Ser Val Glu Thr Glu Ala Asn Gly Gly Leu Val Glu Lys 3220 3225 3230	9696
gag gtg ctg gcc gcc agc gac gca gcc ctt ttg cgc ttc cgg cgc ctg Glu Val Leu Ala Ala Ser Asp Ala Ala Leu Leu Arg Phe Arg Arg Leu 3235 3240 3245	9744
ctg gtg gct gag ctg cag cgt gcc ttc ttt gac aag cac atc tgg ctg Leu Val Ala Glu Leu Gln Arg Gly Phe Phe Asp Lys His Ile Trp Leu 3250 3255 3260	9792
tcc ata tgg gac cgg ccg cct cgt agc cgt ttc act cgc atc cag agg Ser Ile Trp Asp Arg Pro Pro Arg Ser Arg Phe Thr Arg Ile Gln Arg 3265 3270 3275 3280	9840
gcc acc tgc tgc gtt ctg ctg atc tgc ctg ttc ctg gcc gcc aac gcc Ala Thr Cys Cys Val Leu Leu Ile Cys Leu Phe Leu Gly Ala Asn Ala 3285 3290 3295	9888
gtg tgg tac ggg gct gtt ggc gac tct gcc tac agc acg ggg cat gtg Val Trp Tyr Gly Ala Val Gly Asp Ser Ala Tyr Ser Thr Gly His Val 3300 3305 3310	9936
tcc agg ctg agc ccg ctg agc gtc gac aca gtc gct gtt gcc ctg gtg Ser Arg Leu Ser Pro Leu Ser Val Asp Thr Val Ala Val Gly Leu Val 3315 3320 3325	9984
tcc agc gtg gtt gtc tat ccc gtc tac ctg gcc atc ctt ttt ctg ttc Ser Ser Val Val Val Tyr Pro Val Tyr Leu Ala Ile Leu Phe Leu Phe 3330 3335 3340	10032
cgg atg tcc cgg agc aag gtg gct ggg agc ccg agc ccc aca cct gcc Arg Met Ser Arg Ser Lys Val Ala Gly Ser Pro Ser Pro Thr Pro Ala 3345 3350 3355 3360	10080

3345	3350	3355	3360	
ggg cag cag gtg ctg gac atc gac agc tgc ctg gac tgc tcc gtg ctg				10128
Gly Gln Gln Val Leu Asp Ile Asp Ser Cys Leu Asp Ser Ser Val Leu				
	3365	3370	3375	
gac agc tcc ttc ctg acg ttc tca ggc ctg cag gct gag cag gcc ttc				10176
Asp Ser Ser Phe Leu Thr Phe Ser Gly Leu His Ala Glu Gln Ala Phe				
	3380	3385	3390	
gtt gga cag atg aag agt gac ttg ttt ctg gat gat tct aag agt ctg				10224
Val Gly Gln Met Lys Ser Asp Leu Phe Leu Asp Asp Ser Lys Ser Leu				
	3395	3400	3405	
gtg tgc tgg ccc tcc ggc gag gga acg ctg agt tgg ccg gac ctg ctg				10272
Val Cys Trp Pro Ser Gly Glu Gly Thr Leu Ser Trp Pro Asp Leu Leu				
	3410	3415	3420	
agt gac ccg tcc att gtg ggt agc aat ctg ccg cag ctg gca ccg ggc				10320
Ser Asp Pro Ser Ile Val Gly Ser Asn Leu Arg Gln Leu Ala Arg Gly				
	3425	3430	3435	3440
cag gcg ggc cat ggg ctg ggc cca gag gag gac ggc ttc tcc ctg gcc				10368
Gln Ala Gly His Gly Leu Gly Pro Glu Glu Asp Gly Phe Ser Leu Ala				
	3445	3450	3455	
agc ccc tac tgc cct gcc aaa tcc ttc tca gca tca gat gaa gac ctg				10416
Ser Pro Tyr Ser Pro Ala Lys Ser Phe Ser Ala Ser Asp Glu Asp Leu				
	3460	3465	3470	
atc cag cag gtc ctt gcc gag ggg gtc agc agc cca gcc cct acc caa				10464
Ile Gln Gln Val Leu Ala Glu Gly Val Ser Ser Pro Ala Pro Thr Gln				
	3475	3480	3485	
gac acc cag atg gaa acg gac ctg ctg agc agc ctg tcc agc act cct				10512
Asp Thr His Met Glu Thr Asp Leu Leu Ser Ser Leu Ser Ser Thr Pro				
	3490	3495	3500	
ggg gag aag aca gag acg ctg gcg ctg cag agg ctg ggg gag ctg ggg				10560
Gly Glu Lys Thr Glu Thr Leu Ala Leu Gln Arg Leu Gly Glu Leu Gly				
	3505	3510	3515	3520
cca ccc agc cca ggc ctg aac tgg gaa cag ccc cag gca gcg agg ctg				10608
Pro Pro Ser Pro Gly Leu Asn Trp Glu Gln Pro Gln Ala Ala Arg Leu				
	3525	3530	3535	
tcc agg aca gga ctg gtg gag ggt ctg ccg aag cgc ctg ctg ccg gcc				10656
Ser Arg Thr Gly Leu Val Glu Gly Leu Arg Lys Arg Leu Leu Pro Ala				
	3540	3545	3550	
tgg tgt gcc tcc ctg gcc cac ggg ctg agc ctg ctg ctg gtg gct gtg				10704
Trp Cys Ala Ser Leu Ala His Gly Leu Ser Leu Leu Val Ala Val				
	3555	3560	3565	
gct gtg gct gtc tca ggg tgg gtg ggt gcg agc ttc ccc ccg ggc gtg				10752
Ala Val Ala Val Ser Gly Trp Val Gly Ala Ser Phe Pro Pro Gly Val				
	3570	3575	3580	
agt gtt gcg tgg ctg ctg tcc agc agc gcc agc ttc ctg gcc tca ttc				10800
Ser Val Ala Trp Leu Leu Ser Ser Ser Ala Ser Phe Leu Ala Ser Phe				
	3585	3590	3595	3600
ctc ggc tgg gag cca ctg aag gtc ttg ctg gaa gcc ctg tac ttc tca				10848
Leu Gly Trp Glu Pro Leu Lys Val Leu Leu Glu Ala Leu Tyr Phe Ser				
	3605	3610	3615	
ctg gtg gcc aag ccg ctg cac ccg gat gaa gat gac acc ctg gta gag				10896

Leu Val Ala Lys Arg Leu His Pro Asp Glu Asp Asp Thr Leu Val Glu	
3620 3625 3630	
agg ccg ggt gtg agc cct gtg agc gca cgt gtg ccc cgc gta cgg cca	10944
Ser Pro Ala Val Thr Pro Val Ser Ala Arg Val Pro Arg Val Arg Pro	
3635 3640 3645	
ccc cac ggc ttt gca ctc ttc ctg gcc aag gaa gaa gcc cgc aag gtc	10992
Pro His Gly Phe Ala Leu Phe Leu Ala Lys Glu Glu Ala Arg Lys Val	
3650 3655 3660	
aag agg cta cat ggc atg ctg cgg agc ctc ctg gtg tac atg ctt ttt	11040
Lys Arg Leu His Gly Met Leu Arg Ser Leu Leu Val Tyr Met Leu Phe	
3665 3670 3675 3680	
ctg ctg gtg acc ctg ctg gcc agc tat ggg gat gcc tca tgc cat ggg	11088
Leu Leu Val Thr Leu Leu Ala Ser Tyr Gly Asp Ala Ser Cys His Gly	
3685 3690 3695	
cac gcc tac cgt ctg caa agc gcc atc aag cag gag ctg cac agc cgg	11136
His Ala Tyr Arg Leu Gln Ser Ala Ile Lys Gln Glu Leu His Ser Arg	
3700 3705 3710	
gcc ttc ctg gcc atc aag cgg tct gag gag ctc tgg cca tgg atg gcc	11184
Ala Phe Leu Ala Ile Thr Arg Ser Glu Glu Leu Trp Pro Trp Met Ala	
3715 3720 3725	
cac gtg ctg ctg ccc tac gtc cac ggg aac cag tcc agc cca gag ctg	11232
His Val Leu Leu Pro Tyr Val His Gly Asn Gln Ser Ser Pro Glu Leu	
3730 3735 3740	
ggg ccc cca cgg ctg cgg cag gtg cgg ctg cag gaa gca ctc tac cca	11280
Gly Pro Pro Arg Leu Arg Gln Val Arg Leu Gln Glu Ala Leu Tyr Pro	
3745 3750 3755 3760	
gac cct ccc ggc ccc agc gtc cac acg tgc tgg gcc gca gga ggc ttc	11328
Asp Pro Pro Gly Pro Arg Val His Thr Cys Ser Ala Ala Gly Gly Phe	
3765 3770 3775	
agc acc agc gat tac gac gtt ggc tgg gag agt cct cac aat gcc tgc	11376
Ser Thr Ser Asp Tyr Asp Val Gly Trp Glu Ser Pro His Asn Gly Ser	
3780 3785 3790	
ggg acg tgg gcc tat tca gcc ccg gat ctg ctg ggg gca tgg tcc tgg	11424
Gly Thr Trp Ala Tyr Ser Ala Pro Asp Leu Leu Gly Ala Trp Ser Trp	
3795 3800 3805	
ggc tcc tgt gcc gtg tat gac agc ggg ggc tac gtg cag gag ctg gcc	11472
Gly Ser Cys Ala Val Tyr Asp Ser Gly Gly Tyr Val Gln Glu Leu Gly	
3810 3815 3820	
ctg agc ctg gag gag agc cgc gac cgg ctg cgc ttc ctg cag ctg cac	11520
Leu Ser Leu Glu Glu Ser Arg Asp Arg Leu Arg Phe Leu Gln Leu His	
3825 3830 3835 3840	
aac tgg ctg gac aac agg agc cgc gct gtg ttc ctg gag ctc acg cgc	11568
Asn Trp Leu Asp Asn Arg Ser Arg Ala Val Phe Leu Glu Leu Thr Arg	
3845 3850 3855	
tac agc ccg gcc gtg ggg ctg cac gcc gcc gtc acg ctg cgc ctc gag	11616
Tyr Ser Pro Ala Val Gly Leu His Ala Ala Val Thr Leu Arg Leu Glu	
3860 3865 3870	
ttc ccg gcg gcc ggc cgc gcc ctg gcc gcc ctc agc gtc cgc ccc ttt	11664
Phe Pro Ala Ala Gly Arg Ala Leu Ala Ala Leu Ser Val Arg Pro Phe	
3875 3880 3885	

ggg ctg cgc cgc ctc agc ggc ggc ctc tag ctg cct ctg ctc acc tag Ala Leu Arg Arg Leu Ser Ala Gly Leu Ser Leu Pro Leu Leu Thr Ser 3390 3895 3900	11712
gtg tgc ctg ctg ctg ttc gcc gtg cac ttc gcc gtg gcc gag gcc cgt Val Cys Leu Leu Leu Phe Ala Val His Phe Ala Val Ala Glu Ala Arg 3905 3910 3915 3920	11760
act tgg cac agg gaa ggg cgc tgg cgc gtg ctg cgg ctc gga gcc tgg Thr Trp His Arg Glu Gly Arg Trp Arg Val Leu Arg Leu Gly Ala Trp 3925 3930 3935	11803
ggg cgg tgg ctg ctg gtg ggc ctg acg ggc gcc acg gca ctg gta cgc Ala Arg Trp Leu Leu Val Ala Leu Thr Ala Ala Thr Ala Leu Val Arg 3940 3945 3950	11856
ctc gcc cag ctg ggt gcc gct gac cgc cag tgg acc cgt ttc gtg cgc Leu Ala Gln Leu Gly Ala Ala Asp Arg Gln Trp Thr Arg Phe Val Arg 3955 3960 3965	11904
ggc cgc cgc cgc cgc ttc act agc ttc gac cag gtg ggc cac gtg acc Gly Arg Pro Arg Arg Phe Thr Ser Phe Asp Gln Val Ala His Val Ser 3970 3975 3980	11952
tcg gca gcc cgt ggc ctg ggc gcc tag ctg ctc ttc ctg cct ttg gtc Ser Ala Ala Arg Gly Leu Ala Ala Ser Leu Leu Phe Leu Leu Leu Val 3985 3990 3995 4000	12000
aag ggt gcc cag cac gta cgc ttc gtg cgc cag tgg tcg gtc ttt gcc Lys Ala Ala Gln His Val Arg Phe Val Arg Gln Trp Ser Val Phe Gly 4005 4010 4015	12048
aag aca tta tgc cga gct ctg cca gag ctc ctg ggc gtc acc ttg gcc Lys Thr Leu Cys Arg Ala Leu Pro Glu Leu Leu Gly Val Thr Leu Gly 4020 4025 4030	12096
ctg gtg gtg ctc ggc gta gcc tac gcc cag ctg gcc acg ctg ctc gtg Leu Val Val Leu Gly Val Ala Tyr Ala Gln Leu Ala Ile Leu Leu Val 4035 4040 4045	12144
tct tcc tgt gtg gac tcc ctc tgg agc gtg gcc cag gcc ctg ttg gtg Ser Ser Cys Val Asp Ser Leu Trp Ser Val Ala Gln Ala Leu Leu Val 4050 4055 4060	12192
ctg tgc cct ggc act ggc ctc tct acc ctg tgt cct gcc gag tcc tgg Leu Cys Pro Gly Thr Gly Leu Ser Thr Leu Cys Pro Ala Glu Ser Trp 4065 4070 4075 4080	12240
cac ctg tca ccc ctg ctg tgt gtg ggc ctc tgg gca ctg cgg ctg tgg His Leu Ser Pro Leu Leu Cys Val Gly Leu Trp Ala Leu Arg Leu Trp 4085 4090 4095	12288
ggc gcc cta cgg ctg ggc gct gtt att ctc cgc tgg cgc tac cac gcc Gly Ala Leu Arg Leu Gly Ala Val Ile Leu Arg Trp Arg Tyr His Ala 4100 4105 4110	12336
ttg cgt gga gag ctg tac cgg ccg gcc tgg gag ccc cag gac tac gag Leu Arg Gly Glu Leu Tyr Arg Pro Ala Trp Glu Pro Gln Asp Tyr Glu 4115 4120 4125	12384
atg gtg gag ttg ttc ctg cgc agg ctg cgc ctc tgg atg ggc ctc agc Met Val Glu Leu Phe Leu Arg Arg Leu Arg Leu Trp Met Gly Leu Ser 4130 4135 4140	12432
aag gtc aag gag ttc cgc cac aaa gtc cgc ttt gaa ggc atg gag ccg Lys Val Lys Glu Phe Arg His Lys Val Arg Phe Glu Gly Met Glu Pro 4145 4150 4155 4160	12480

ctg ccc tct cgg tcc tcc agg ggc tcc aag gta tcc cgg gat gtg ccc 12528
 Leu Pro Ser Arg Ser Ser Arg Gly Ser Lys Val Ser Pro Asp Val Pro
 4155 4170 4175
 cca ccc agc ggt ggc tcc gat gcc tgg cac ccc tcc acc tcc tcc agc 12576
 Pro Pro Ser Ala Gly Ser Asp Ala Ser His Pro Ser Thr Ser Ser Ser
 4180 4185 4190
 cag ctg gat ggg ctg agc gtg agc ctg ggc cgg ctg ggg aca agg tgt 12624
 Gln Leu Asp Gly Leu Ser Val Ser Leu Gly Arg Leu Gly Thr Arg Cys
 4195 4200 4205
 gag cct gag ccc tcc cgg ctc caa gcc gtg ttc gag gcc ctg ctc acc 12672
 Glu Pro Glu Pro Ser Arg Leu Gln Ala Val Phe Glu Ala Leu Leu Thr
 4210 4215 4220
 cag ttt gac cga ctc aac cag gcc aca gag gac gtc tac cag ctg gag 12720
 Gln Phe Asp Arg Leu Asn Gln Ala Thr Glu Asp Val Tyr Gln Leu Glu
 4225 4230 4235 4240
 cag cag ctg cac agc ctg caa ggc cgg agc agc agc cgg ggc ccc gcc 12768
 Gln Gln Leu His Ser Leu Gln Gly Arg Arg Ser Ser Arg Ala Pro Ala
 4245 4250 4255
 gga tct tcc cgt ggc cca tcc cgg ggc ctg cgg cca gca ctg ccc agc 12816
 Gly Ser Ser Arg Gly Pro Ser Pro Gly Leu Arg Pro Ala Leu Pro Ser
 4260 4265 4270
 cgc ctt gcc cgg gcc agt cgg ggt gtg gac ctg gcc act ggc ccc agc 12864
 Arg Leu Ala Arg Ala Ser Arg Gly Val Asp Leu Ala Thr Gly Pro Ser
 4275 4280 4285
 agg aca ccc ctt cgg gcc aag aac aag gtc cac ccc agc agc act tag 12912
 Arg Thr Pro Leu Arg Ala Lys Asn Lys Val His Pro Ser Ser Thr
 4290 4295 4300

<210> 2
 <211> 4303
 <212> PRT
 <213> Homo sapiens PKD-1 protein

<400> 2
 Met Pro Pro Ala Ala Pro Ala Arg Leu Ala Leu Ala Leu Gly Leu Gly
 1 5 10 15
 Leu Trp Leu Gly Ala Leu Ala Gly Gly Pro Gly Arg Gly Cys Gly Pro
 20 25 30
 Cys Glu Pro Pro Cys Leu Cys Gly Pro Ala Pro Gly Ala Ala Cys Arg
 35 40 45
 Val Asn Cys Ser Gly Arg Gly Leu Arg Thr Leu Gly Pro Ala Leu Arg
 50 55 60
 Ile Pro Ala Asp Ala Thr Glu Leu Asp Val Ser His Asn Leu Leu Arg
 65 70 75 80
 Ala Leu Asp Val Gly Leu Leu Ala Asn Leu Ser Ala Leu Ala Glu Leu
 85 90 95
 Asp Ile Ser Asn Asn Lys Ile Ser Thr Leu Glu Glu Gly Ile Phe Ala
 100 105 110
 Asn Leu Phe Asn Leu Ser Glu Ile Asn Leu Ser Gly Asn Pro Phe Glu
 115 120 125

Cys	Asp	Cys	Gly	Leu	Ala	Trp	Leu	Pro	Gln	Trp	Ala	Glu	Glu	Gln	Gln	130	135	140
Val	Arg	Val	Val	Gln	Pro	Glu	Ala	Ala	Thr	Cys	Ala	Gly	Pro	Gly	Ser	145	150	155
Leu	Ala	Gly	Gln	Pro	Leu	Leu	Gly	Ile	Pro	Leu	Leu	Asp	Ser	Gly	Cys	165	170	175
Gly	Glu	Glu	Tyr	Val	Ala	Cys	Leu	Pro	Asp	Asn	Ser	Ser	Gly	Thr	Val	190	185	190
Ala	Ala	Val	Ser	Phe	Ser	Ala	Ala	His	Glu	Gly	Leu	Leu	Gln	Pro	Glu	195	200	205
Ala	Cys	Ser	Ala	Phe	Cys	Phe	Ser	Thr	Gly	Gln	Gly	Leu	Ala	Ala	Leu	210	215	220
Ser	Glu	Gln	Gly	Trp	Cys	Leu	Cys	Gly	Ala	Ala	Gln	Pro	Ser	Ser	Ala	225	230	235
Ser	Phe	Ala	Cys	Leu	Ser	Leu	Cys	Ser	Gly	Pro	Pro	Ala	Pro	Pro	Ala	245	250	255
Pro	Thr	Cys	Arg	Gly	Pro	Thr	Leu	Leu	Gln	His	Val	Phe	Pro	Ala	Ser	260	265	270
Pro	Gly	Ala	Thr	Leu	Val	Gly	Pro	His	Gly	Pro	Leu	Ala	Ser	Gly	Gln	275	280	285
Leu	Ala	Ala	Phe	His	Ile	Ala	Ala	Pro	Leu	Pro	Val	Thr	Asp	Thr	Arg	290	295	300
Trp	Asp	Phe	Gly	Asp	Gly	Ser	Ala	Glu	Val	Asp	Ala	Ala	Gly	Pro	Ala	305	310	315
Ala	Ser	His	Arg	Tyr	Val	Leu	Pro	Gly	Arg	Tyr	His	Val	Thr	Ala	Val	325	330	335
Leu	Ala	Leu	Gly	Ala	Gly	Ser	Ala	Leu	Leu	Gly	Thr	Asp	Val	Gln	Val	340	345	350
Glu	Ala	Ala	Pro	Ala	Ala	Leu	Glu	Leu	Val	Cys	Pro	Ser	Ser	Val	Gln	355	360	365
Ser	Asp	Glu	Ser	Leu	Asp	Leu	Ser	Ile	Gln	Asn	Arg	Gly	Gly	Ser	Gly	370	375	380
Leu	Glu	Ala	Ala	Tyr	Ser	Ile	Val	Ala	Leu	Gly	Glu	Glu	Pro	Ala	Arg	385	390	395
Ala	Val	His	Pro	Leu	Cys	Pro	Ser	Asp	Thr	Glu	Ile	Phe	Pro	Gly	Asn	405	410	415
Gly	His	Cys	Tyr	Arg	Leu	Val	Val	Glu	Lys	Ala	Ala	Trp	Leu	Gln	Ala	420	425	430
Gln	Glu	Gln	Cys	Gln	Ala	Trp	Ala	Gly	Ala	Ala	Leu	Ala	Met	Val	Asp	435	440	445
Ser	Pro	Ala	Val	Gln	Arg	Phe	Leu	Val	Ser	Arg	Val	Thr	Arg	Ser	Leu	450	455	460
Asp	Val	Trp	Ile	Gly	Phe	Ser	Thr	Val	Gln	Gly	Val	Glu	Val	Gly	Pro	465	470	475

Ala Pro Gln Gly Glu Ala Phe Ser Leu Glu Ser Cys Gln Asn Trp Leu
 485 490 495
 Pro Gly Glu Pro His Pro Ala Thr Ala Glu His Cys Val Arg Leu Gly
 500 505 510
 Pro Thr Gly Trp Cys Asn Thr Asp Leu Cys Ser Ala Pro His Ser Tyr
 515 520 525
 Val Cys Glu Leu Gln Pro Gly Gly Pro Val Gln Asp Ala Glu Asn Leu
 530 535 540
 Leu Val Gly Ala Pro Ser Gly Asp Leu Gln Gly Pro Leu Thr Pro Leu
 545 550 555
 Ala Gln Gln Asp Gly Leu Ser Ala Pro His Glu Pro Val Glu Val Met
 565 570 575
 Val Phe Pro Gly Leu Arg Leu Ser Arg Glu Ala Phe Leu Thr Thr Ala
 580 585 590
 Glu Phe Gly Thr Gln Glu Leu Arg Arg Pro Ala Gln Leu Arg Leu Gln
 595 600 605
 Val Tyr Arg Leu Leu Ser Thr Ala Gly Thr Pro Glu Asn Gly Ser Glu
 610 615 620
 Pro Glu Ser Arg Ser Pro Asp Asn Arg Thr Gln Leu Ala Pro Ala Cys
 625 630 635 640
 Met Pro Gly Gly Arg Trp Cys Pro Gly Ala Asn Ile Cys Leu Pro Leu
 645 650 655
 Asp Ala Ser Cys His Pro Gln Ala Cys Ala Asn Gly Cys Thr Ser Gly
 660 665 670
 Pro Gly Leu Pro Gly Ala Pro Tyr Ala Leu Trp Arg Glu Phe Leu Phe
 675 680 685
 Ser Val Pro Ala Gly Pro Pro Ala Gln Tyr Ser Val Thr Leu His Gly
 690 695 700
 Gln Asp Val Leu Met Leu Pro Gly Asp Leu Val Gly Leu Gln His Asp
 705 710 715 720
 Ala Gly Pro Gly Ala Leu Leu His Cys Ser Pro Ala Pro Gly His Pro
 725 730 735
 Gly Pro Arg Ala Pro Tyr Leu Ser Ala Asn Ala Ser Ser Trp Leu Pro
 740 745 750
 His Leu Pro Ala Gln Leu Glu Gly Thr Trp Gly Cys Pro Ala Cys Ala
 755 760 765
 Leu Arg Leu Leu Ala Gln Arg Glu Gln Leu Thr Val Leu Leu Gly Leu
 770 775 780
 Arg Pro Asn Pro Gly Leu Arg Leu Pro Gly Arg Tyr Glu Val Arg Ala
 785 790 795 800
 Glu Val Gly Asn Gly Val Ser Arg His Asn Leu Ser Cys Ser Phe Asp
 805 810 815
 Val Val Ser Pro Val Ala Gly Leu Arg Val Ile Tyr Pro Ala Pro Arg
 820 825 830
 Asp Gly Arg Leu Tyr Val Pro Thr Asn Gly Ser Ala Leu Val Leu Gln

835		840		845
Val Asp Ser Gly Ala Asn Ala Thr Ala Thr Ala Arg Trp Pro Gly Gly	850	855	860	
Ser Leu Ser Ala Arg Phe Glu Asn Val Cys Pro Ala Leu Val Ala Thr	865	870	875	880
Phe Val Pro Ala Cys Pro Trp Glu Thr Asn Asp Thr Leu Phe Ser Val	885	890	895	
Val Ala Leu Pro Trp Leu Ser Glu Gly Glu His Val Val Asp Val Val	900	905	910	
Val Glu Asn Ser Ala Ser Arg Ala Asn Leu Ser Leu Arg Val Thr Ala	915	920	925	
Glu Glu Pro Ile Cys Gly Leu Arg Ala Thr Pro Ser Pro Glu Ala Arg	930	935	940	
Val Leu Gln Gly Val Leu Val Arg Tyr Ser Pro Val Val Glu Ala Gly	945	950	955	960
Ser Asp Met Val Phe Arg Trp Thr Ile Asn Asp Lys Gln Ser Leu Thr	965	970	975	
Phe Gln Asn Val Val Phe Asn Val Ile Tyr Gln Ser Ala Ala Val Phe	980	985	990	
Lys Leu Ser Leu Thr Ala Ser Asn His Val Ser Asn Val Thr Val Asn	995	1000	1005	
Tyr Asn Val Thr Val Glu Arg Met Asn Arg Met Gln Gly Leu Gln Val	1010	1015	1020	
Ser Thr Val Pro Ala Val Leu Ser Pro Asn Ala Thr Leu Ala Leu Thr	1025	1030	1035	1040
Ala Gly Val Leu Val Asp Ser Ala Val Glu Val Ala Phe Leu Trp Thr	1045	1050	1055	
Phe Gly Asp Gly Glu Gln Ala Leu His Gln Phe Gln Pro Pro Tyr Asn	1060	1065	1070	
Glu Ser Phe Pro Val Pro Asp Pro Ser Val Ala Gln Val Leu Val Glu	1075	1080	1085	
His Asn Val Thr His Thr Tyr Ala Ala Pro Gly Glu Tyr Leu Leu Thr	1090	1095	1100	
Val Leu Ala Ser Asn Ala Phe Glu Asn Leu Thr Gln Gln Val Pro Val	1105	1110	1115	1120
Ser Val Arg Ala Ser Leu Pro Ser Val Ala Val Gly Val Ser Asp Gly	1125	1130	1135	
Val Leu Val Ala Gly Arg Pro Val Thr Phe Tyr Pro His Pro Leu Pro	1140	1145	1150	
Ser Pro Gly Gly Val Leu Tyr Thr Trp Asp Phe Gly Asp Gly Ser Pro	1155	1160	1165	
Val Leu Thr Gln Ser Gln Pro Ala Ala Asn His Thr Tyr Ala Ser Arg	1170	1175	1180	
Gly Thr Tyr His Val Arg Leu Glu Val Asn Asn Thr Val Ser Gly Ala	1185	1190	1195	1200

Ala Ala Gln Ala Asp Val Arg Val Phe Glu Glu Leu Arg Gly Leu Ser
 1205 1210 1215
 Val Asp Met Ser Leu Ala Val Glu Gln Gly Ala Pro Val Val Val Ser
 1220 1225 1230
 Ala Ala Val Gln Thr Gly Asp Asn Ile Thr Trp Thr Phe Asp Met Gly
 1235 1240 1245
 Asp Gly Thr Val Leu Ser Gly Pro Glu Ala Thr Val Glu His Val Tyr
 1250 1255 1260
 Leu Arg Ala Gln Asn Cys Thr Val Thr Val Gly Ala Gly Ser Pro Ala
 1265 1270 1275 1280
 Gly His Leu Ala Arg Ser Leu His Val Leu Val Phe Val Leu Glu Val
 1285 1290 1295
 Leu Arg Val Glu Pro Ala Ala Cys Ile Pro Thr Gln Pro Asp Ala Arg
 1300 1305 1310
 Leu Thr Ala Tyr Val Thr Gly Asn Pro Ala His Tyr Leu Phe Asp Trp
 1315 1320 1325
 Thr Phe Gly Asp Gly Ser Ser Asn Thr Thr Val Arg Gly Cys Pro Thr
 1330 1335 1340
 Val Thr His Asn Phe Thr Arg Ser Gly Thr Phe Pro Leu Ala Leu Val
 1345 1350 1355 1360
 Leu Ser Ser Arg Val Asn Arg Ala His Tyr Phe Thr Ser Ile Cys Val
 1365 1370 1375
 Glu Pro Glu Val Gly Asn Val Thr Leu Gln Pro Glu Arg Gln Phe Val
 1380 1385 1390
 Gln Leu Gly Asp Glu Ala Trp Leu Val Ala Cys Ala Trp Pro Pro Phe
 1395 1400 1405
 Pro Tyr Arg Tyr Thr Trp Asp Phe Gly Thr Glu Glu Ala Ala Pro Thr
 1410 1415 1420
 Arg Ala Arg Gly Pro Glu Val Thr Phe Ile Tyr Arg Asp Pro Gly Ser
 1425 1430 1435 1440
 Tyr Leu Val Thr Val Thr Ala Ser Asn Asn Ile Ser Ala Ala Asn Asp
 1445 1450 1455
 Ser Ala Leu Val Glu Val Gln Glu Pro Val Leu Val Thr Ser Ile Lys
 1460 1465 1470
 Val Asn Gly Ser Leu Gly Leu Glu Leu Gln Gln Pro Tyr Leu Phe Ser
 1475 1480 1485
 Ala Val Gly Arg Gly Arg Pro Ala Ser Tyr Leu Trp Asp Leu Gly Asp
 1490 1495 1500
 Gly Gly Trp Leu Glu Gly Pro Glu Val Thr His Ala Tyr Asn Ser Thr
 1505 1510 1515 1520
 Gly Asp Phe Thr Val Arg Val Ala Gly Trp Asn Glu Val Ser Arg Ser
 1525 1530 1535
 Glu Ala Trp Leu Asn Val Thr Val Lys Arg Arg Val Arg Gly Leu Val
 1540 1545 1550

Val Asn Ala Ser Arg Thr Val Val Pro Leu Asn Gly Ser Val Ser Phe
 1555 1560 1565
 Ser Thr Ser Leu Glu Ala Gly Ser Asp Val Arg Tyr Ser Trp Val Leu
 1570 1575 1580
 Cys Asp Arg Cys Thr Pro Ile Pro Gly Gly Pro Thr Ile Ser Tyr Thr
 535 1590 1595 1600
 Phe Arg Ser Val Gly Thr Phe Asn Ile Ile Val Thr Ala Glu Asn Glu
 1605 1610 1615
 Val Gly Ser Ala Gln Asp Ser Ile Phe Val Tyr Val Leu Gln Leu Ile
 1620 1625 1630
 Glu Gly Leu Gln Val Val Gly Gly Gly Arg Tyr Phe Pro Thr Asn His
 1635 1640 1645
 Thr Val Gln Leu Gln Ala Val Val Arg Asp Gly Thr Asn Val Ser Tyr
 1650 1655 1660
 Ser Trp Thr Ala Trp Arg Asp Arg Gly Pro Ala Leu Ala Gly Ser Gly
 665 1670 1675 1680
 Lys Gly Phe Ser Leu Thr Val Leu Glu Ala Gly Thr Tyr His Val Gln
 1685 1690 1695
 Leu Arg Ala Thr Asn Met Leu Gly Ser Ala Trp Ala Asp Cys Thr Met
 1700 1705 1710
 Asp Phe Val Glu Pro Val Gly Trp Leu Met Val Ala Ala Ser Pro Asn
 1715 1720 1725
 Pro Ala Ala Val Asn Thr Ser Val Thr Leu Ser Ala Glu Leu Ala Gly
 1730 1735 1740
 Gly Ser Gly Val Val Tyr Thr Trp Ser Leu Glu Glu Gly Leu Ser Trp
 745 1750 1755 1760
 Glu Thr Ser Glu Pro Phe Thr Thr His Ser Phe Pro Thr Pro Gly Leu
 1765 1770 1775
 His Leu Val Thr Met Thr Ala Gly Asn Pro Leu Gly Ser Ala Asn Ala
 1780 1785 1790
 Thr Val Glu Val Asp Val Gln Val Pro Val Ser Gly Leu Ser Ile Arg
 1795 1800 1805
 Ala Ser Glu Pro Gly Gly Ser Phe Val Ala Ala Gly Ser Ser Val Pro
 1810 1815 1820
 Phe Trp Gly Gln Leu Ala Thr Gly Thr Asn Val Ser Trp Cys Trp Ala
 825 1830 1835 1840
 Val Pro Gly Gly Ser Ser Lys Arg Gly Pro His Val Thr Met Val Phe
 1845 1850 1855
 Pro Asp Ala Gly Thr Phe Ser Ile Arg Leu Asn Ala Ser Asn Ala Val
 1860 1865 1870
 Ser Trp Val Ser Ala Thr Tyr Asn Leu Thr Ala Glu Glu Pro Ile Val
 1875 1880 1885
 Gly Leu Val Leu Trp Ala Ser Ser Lys Val Val Ala Pro Gly Gln Leu
 1890 1895 1900
 Val His Phe Gln Ile Leu Leu Ala Ala Gly Ser Ala Val Thr Phe Arg

1905	1910	1915	1920
Leu Gln Val Gly Gly Ala Asn Pro Glu Val Leu Pro Gly Pro Arg Phe 1925 1930 1935			
Ser His Ser Phe Pro Arg Val Gly Asp His Val Val Ser Val Arg Gly 1940 1945 1950			
Lys Asn His Val Ser Trp Ala Gln Ala Gln Val Arg Ile Val Val Leu 1955 1960 1965			
Glu Ala Val Ser Gly Leu Gln Val Pro Asn Cys Cys Glu Pro Gly Ile 1970 1975 1980			
Ala Thr Gly Thr Glu Arg Asn Phe Thr Ala Arg Val Gln Arg Gly Ser 1985 1990 1995 2000			
Arg Val Ala Tyr Ala Trp Tyr Phe Ser Leu Gln Lys Val Gln Gly Asp 2005 2010 2015			
Ser Leu Val Ile Leu Ser Gly Arg Asp Val Thr Tyr Thr Pro Val Ala 2020 2025 2030			
Ala Gly Leu Leu Glu Ile Gln Val Arg Ala Phe Asn Ala Leu Gly Ser 2035 2040 2045			
Glu Asn Arg Thr Leu Val Leu Glu Val Gln Asp Ala Val Gln Tyr Val 2050 2055 2060			
Ala Leu Gln Ser Gly Pro Cys Phe Thr Asn Arg Ser Ala Gln Phe Glu 2065 2070 2075 2080			
Ala Ala Thr Ser Pro Ser Pro Arg Arg Val Ala Tyr His Trp Asp Phe 2085 2090 2095			
Gly Asp Gly Ser Pro Gly Gln Asp Thr Asp Glu Pro Arg Ala Glu His 2100 2105 2110			
Ser Tyr Leu Arg Pro Gly Asp Tyr Arg Val Gln Val Asn Ala Ser Asn 2115 2120 2125			
Leu Val Ser Phe Phe Val Ala Gln Ala Thr Val Thr Val Gln Val Leu 2130 2135 2140			
Ala Cys Arg Glu Pro Glu Val Asp Val Val Leu Pro Leu Gln Val Leu 2145 2150 2155 2160			
Met Arg Arg Ser Gln Arg Asn Tyr Leu Glu Ala His Val Asp Leu Arg 2165 2170 2175			
Asp Cys Val Thr Tyr Gln Thr Glu Tyr Arg Trp Glu Val Tyr Arg Thr 2180 2185 2190			
Ala Ser Cys Gln Arg Pro Gly Arg Pro Ala Arg Val Ala Leu Pro Gly 2195 2200 2205			
Val Asp Val Ser Arg Pro Arg Leu Val Leu Pro Arg Leu Ala Leu Pro 2210 2215 2220			
Val Gly His Tyr Cys Phe Val Phe Val Val Ser Phe Gly Asp Thr Pro 2225 2230 2235 2240			
Leu Thr Gln Ser Ile Gln Ala Asn Val Thr Val Ala Pro Glu Arg Leu 2245 2250 2255			
Val Pro Ile Ile Glu Gly Gly Ser Tyr Arg Val Trp Ser Asp Thr Arg 2260 2265 2270			

Asp Leu Val Leu Asp Gly Ser Glu Ser Tyr Asp Pro Asn Leu Glu Asp
 2275 2280 2285
 Gly Asp Gln Thr Pro Leu Ser Phe His Trp Ala Cys Val Ala Ser Thr
 2290 2295 2300
 Gln Arg Glu Ala Gly Gly Cys Ala Leu Asn Phe Gly Pro Arg Gly Ser
 305 2310 2315 2320
 Ser Thr Val Thr Ile Pro Arg Glu Arg Leu Ala Ala Gly Val Glu Tyr
 2325 2330 2335
 Thr Phe Ser Leu Thr Val Trp Lys Ala Gly Arg Lys Glu Glu Ala Thr
 2340 2345 2350
 Asn Gln Thr Val Leu Ile Arg Ser Gly Arg Val Pro Ile Val Ser Leu
 2355 2360 2365
 Glu Cys Val Ser Cys Lys Ala Gln Ala Val Tyr Glu Val Ser Arg Ser
 2370 2375 2380
 Ser Tyr Val Tyr Leu Glu Gly Arg Cys Leu Asn Cys Ser Ser Gly Ser
 335 2390 2395 2400
 Lys Arg Gly Arg Trp Ala Ala Arg Thr Phe Ser Asn Lys Thr Leu Val
 2405 2410 2415
 Leu Asp Glu Thr Thr Thr Ser Thr Gly Ser Ala Gly Met Arg Leu Val
 2420 2425 2430
 Leu Arg Arg Gly Val Leu Arg Asp Gly Glu Gly Tyr Thr Phe Thr Leu
 2435 2440 2445
 Thr Val Leu Gly Arg Ser Gly Glu Glu Glu Gly Cys Ala Ser Ile Arg
 2450 2455 2460
 Leu Ser Pro Asn Arg Pro Pro Leu Gly Gly Ser Cys Arg Leu Phe Pro
 455 2470 2475 2480
 Leu Gly Ala Val His Ala Leu Thr Thr Lys Val His Phe Glu Cys Thr
 2485 2490 2495
 Gly Trp His Asp Ala Glu Asp Ala Gly Ala Pro Leu Val Tyr Ala Leu
 2500 2505 2510
 Leu Leu Arg Arg Cys Arg Gln Gly His Cys Glu Glu Phe Cys Val Tyr
 2515 2520 2525
 Lys Gly Ser Leu Ser Ser Tyr Gly Ala Val Leu Pro Pro Gly Phe Arg
 2530 2535 2540
 Pro His Phe Glu Val Gly Leu Ala Val Val Val Gln Asp Gln Leu Gly
 545 2550 2555 2560
 Ala Ala Val Val Ala Leu Asn Arg Ser Leu Ala Ile Thr Leu Pro Glu
 2565 2570 2575
 Pro Asn Gly Ser Ala Thr Gly Leu Thr Val Trp Leu His Gly Leu Thr
 2580 2585 2590
 Ala Ser Val Leu Pro Gly Leu Leu Arg Gln Ala Asp Pro Gln His Val
 2595 2600 2605
 Ile Glu Tyr Ser Leu Ala Leu Val Thr Val Leu Asn Glu Tyr Glu Arg
 2610 2615 2620

Ala Leu Asp Val Ala Ala Glu Pro Lys His Glu Arg Gln His Arg Ala
 625 2630 2635 2640
 Gln Ile Arg Lys Asn Ile Thr Glu Thr Leu Val Ser Leu Arg Val His
 2645 2650 2655
 Thr Val Asp Asp Ile Gln Gln Ile Ala Ala Ala Leu Ala Gln Cys Met
 2660 2665 2670
 Gly Pro Ser Arg Glu Leu Val Cys Arg Ser Cys Leu Lys Gln Thr Leu
 2675 2680 2685
 His Lys Leu Glu Ala Met Met Leu Ile Leu Gln Ala Glu Thr Thr Ala
 2690 2695 2700
 Gly Thr Val Thr Pro Thr Ala Ile Gly Asp Ser Ile Leu Asn Ile Thr
 705 2710 2715 2720
 Gly Asp Leu Ile His Leu Ala Ser Ser Asp Val Arg Ala Pro Gln Pro
 2725 2730 2735
 Ser Glu Leu Gly Ala Glu Ser Pro Ser Arg Met Val Ala Ser Gln Ala
 2740 2745 2750
 Tyr Asn Leu Thr Ser Ala Leu Met Arg Ile Leu Met Arg Ser Arg Val
 2755 2760 2765
 Leu Asn Glu Glu Pro Leu Thr Leu Ala Gly Glu Glu Ile Val Ala Gln
 2770 2775 2780
 Gly Lys Arg Ser Asp Pro Arg Ser Leu Leu Cys Tyr Gly Gly Ala Pro
 735 2790 2795 2800
 Gly Pro Gly Cys His Phe Ser Ile Pro Glu Ala Phe Ser Gly Ala Leu
 2805 2810 2815
 Ala Asn Leu Ser Asp Val Val Gln Leu Ile Phe Leu Val Asp Ser Asn
 2820 2825 2830
 Pro Phe Pro Phe Gly Tyr Ile Ser Asn Tyr Thr Val Ser Thr Lys Val
 2835 2840 2845
 Ala Ser Met Ala Phe Gln Thr Gln Ala Gly Ala Gln Ile Pro Ile Glu
 2850 2855 2860
 Arg Leu Ala Ser Glu Arg Ala Ile Thr Val Lys Val Pro Asn Asn Ser
 865 2870 2875 2880
 Asp Trp Ala Ala Arg Gly His Arg Ser Ser Ala Asn Ser Ala Asn Ser
 2885 2890 2895
 Val Val Val Gln Pro Gln Ala Ser Val Gly Ala Val Val Thr Leu Asp
 2900 2905 2910
 Ser Ser Asn Pro Ala Ala Gly Leu His Leu Gln Leu Asn Tyr Thr Leu
 2915 2920 2925
 Leu Asp Gly His Tyr Leu Ser Glu Glu Pro Glu Pro Tyr Leu Ala Val
 2930 2935 2940
 Tyr Leu His Ser Glu Pro Arg Pro Asn Glu His Asn Cys Ser Ala Ser
 945 2950 2955 2960
 Arg Arg Ile Arg Pro Glu Ser Leu Gln Gly Ala Asp His Arg Pro Tyr
 2965 2970 2975
 Thr Phe Phe Ile Ser Pro Gly Ser Arg Asp Pro Ala Gly Ser Tyr His

2980

2985

2990

Leu Asn Leu Ser Ser His Phe Arg Trp Ser Ala Leu Gln Val Ser Val
 2995 3000 3005
 Gly Leu Tyr Thr Ser Leu Cys Gln Tyr Phe Ser Glu Glu Asp Met Val
 3010 3015 3020
 Trp Arg Thr Glu Gly Leu Leu Pro Leu Glu Glu Thr Ser Pro Arg Gln
 3025 3030 3035 3040
 Ala Val Cys Leu Thr Arg His Leu Thr Ala Phe Gly Ala Ser Leu Phe
 3045 3050 3055
 Val Pro Pro Ser His Val Arg Phe Val Phe Pro Glu Pro Thr Ala Asp
 3060 3065 3070
 Val Asn Tyr Ile Val Met Leu Thr Cys Ala Val Cys Leu Val Thr Tyr
 3075 3080 3085
 Met Val Met Ala Ala Ile Leu His Lys Leu Asp Gln Leu Asp Ala Ser
 3090 3095 3100
 Arg Gly Arg Ala Ile Pro Phe Cys Gly Gln Arg Gly Arg Phe Lys Tyr
 3105 3110 3115 3120
 Glu Ile Leu Val Lys Thr Gly Trp Gly Arg Gly Ser Gly Thr Thr Ala
 3125 3130 3135
 His Val Gly Ile Met Leu Tyr Gly Val Asp Ser Arg Ser Gly His Arg
 3140 3145 3150
 His Leu Asp Gly Asp Arg Ala Phe His Arg Asn Ser Leu Asp Ile Phe
 3155 3160 3165
 Arg Ile Ala Thr Pro His Ser Leu Gly Ser Val Trp Lys Ile Arg Val
 3170 3175 3180
 Trp His Asp Asn Lys Gly Leu Ser Pro Ala Trp Phe Leu Gln His Val
 3185 3190 3195 3200
 Ile Val Arg Asp Leu Gln Thr Ala Arg Ser Ala Phe Phe Leu Val Asn
 3205 3210 3215
 Asp Trp Leu Ser Val Glu Thr Glu Ala Asn Gly Gly Leu Val Glu Lys
 3220 3225 3230
 Glu Val Leu Ala Ala Ser Asp Ala Ala Leu Leu Arg Phe Arg Arg Leu
 3235 3240 3245
 Leu Val Ala Glu Leu Gln Arg Gly Phe Phe Asp Lys His Ile Trp Leu
 3250 3255 3260
 Ser Ile Trp Asp Arg Pro Pro Arg Ser Arg Phe Thr Arg Ile Gln Arg
 3265 3270 3275 3280
 Ala Thr Cys Cys Val Leu Leu Ile Cys Leu Phe Leu Gly Ala Asn Ala
 3285 3290 3295
 Val Trp Tyr Gly Ala Val Gly Asp Ser Ala Tyr Ser Thr Gly His Val
 3300 3305 3310
 Ser Arg Leu Ser Pro Leu Ser Val Asp Thr Val Ala Val Gly Leu Val
 3315 3320 3325
 Ser Ser Val Val Val Tyr Pro Val Tyr Leu Ala Ile Leu Phe Leu Phe
 3330 3335 3340

Arg Met Ser Arg Ser Lys Val Ala Gly Ser Pro Ser Pro Thr Pro Ala
 345 3350 3355 3360
 Gly Gln Gln Val Leu Asp Ile Asp Ser Cys Leu Asp Ser Ser Val Leu
 3365 3370 3375
 Asp Ser Ser Phe Leu Thr Phe Ser Gly Leu His Ala Glu Gln Ala Phe
 3380 3385 3390
 Val Gly Gln Met Lys Ser Asp Leu Phe Leu Asp Asp Ser Lys Ser Leu
 3395 3400 3405
 Val Cys Trp Pro Ser Gly Glu Gly Thr Leu Ser Trp Pro Asp Leu Leu
 3410 3415 3420
 Ser Asp Pro Ser Ile Val Gly Ser Asn Leu Arg Gln Leu Ala Arg Gly
 425 3430 3435 3440
 Gln Ala Gly His Gly Leu Gly Pro Glu Glu Asp Gly Phe Ser Leu Ala
 3445 3450 3455
 Ser Pro Tyr Ser Pro Ala Lys Ser Phe Ser Ala Ser Asp Glu Asp Leu
 3460 3465 3470
 Ile Gln Gln Val Leu Ala Glu Gly Val Ser Ser Pro Ala Pro Thr Gln
 3475 3480 3485
 Asp Thr His Met Glu Thr Asp Leu Leu Ser Ser Leu Ser Ser Thr Pro
 3490 3495 3500
 Gly Glu Lys Thr Glu Thr Leu Ala Leu Gln Arg Leu Gly Glu Leu Gly
 505 3510 3515 3520
 Pro Pro Ser Pro Gly Leu Asn Trp Glu Gln Pro Gln Ala Ala Arg Leu
 3525 3530 3535
 Ser Arg Thr Gly Leu Val Glu Gly Leu Arg Lys Arg Leu Leu Pro Ala
 3540 3545 3550
 Trp Cys Ala Ser Leu Ala His Gly Leu Ser Leu Leu Leu Val Ala Val
 3555 3560 3565
 Ala Val Ala Val Ser Gly Trp Val Gly Ala Ser Phe Pro Pro Gly Val
 3570 3575 3580
 Ser Val Ala Trp Leu Leu Ser Ser Ser Ala Ser Phe Leu Ala Ser Phe
 585 3590 3595 3600
 Leu Gly Trp Glu Pro Leu Lys Val Leu Leu Glu Ala Leu Tyr Phe Ser
 3605 3610 3615
 Leu Val Ala Lys Arg Leu His Pro Asp Glu Asp Asp Thr Leu Val Glu
 3620 3625 3630
 Ser Pro Ala Val Thr Pro Val Ser Ala Arg Val Pro Arg Val Arg Pro
 3635 3640 3645
 Pro His Gly Phe Ala Leu Phe Leu Ala Lys Glu Glu Ala Arg Lys Val
 3650 3655 3660
 Lys Arg Leu His Gly Met Leu Arg Ser Leu Leu Val Tyr Met Leu Phe
 665 3670 3675 3680
 Leu Leu Val Thr Leu Leu Ala Ser Tyr Gly Asp Ala Ser Cys His Gly
 3685 3690 3695

His Ala Tyr Arg Leu Gln Ser Ala Ile Lys Gln Glu Leu His Ser Arg
 3700 3705 3710
 Ala Phe Leu Ala Ile Thr Arg Ser Glu Glu Leu Trp Pro Trp Met Ala
 3715 3720 3725
 His Val Leu Leu Pro Tyr Val His Gly Asn Gln Ser Ser Pro Glu Leu
 3730 3735 3740
 Gly Pro Pro Arg Leu Arg Gln Val Arg Leu Gln Glu Ala Leu Tyr Pro
 3745 3750 3755 3760
 Asp Pro Pro Gly Pro Arg Val His Thr Cys Ser Ala Ala Gly Gly Phe
 3765 3770 3775
 Ser Thr Ser Asp Tyr Asp Val Gly Trp Glu Ser Pro His Asn Gly Ser
 3780 3785 3790
 Gly Thr Trp Ala Tyr Ser Ala Pro Asp Leu Leu Gly Ala Trp Ser Trp
 3795 3800 3805
 Gly Ser Cys Ala Val Tyr Asp Ser Gly Gly Tyr Val Gln Glu Leu Gly
 3810 3815 3820
 Leu Ser Leu Glu Glu Ser Arg Asp Arg Leu Arg Phe Leu Gln Leu His
 3825 3830 3835 3840
 Asn Trp Leu Asp Asn Arg Ser Arg Ala Val Phe Leu Glu Leu Thr Arg
 3845 3850 3855
 Tyr Ser Pro Ala Val Gly Leu His Ala Ala Val Thr Leu Arg Leu Glu
 3860 3865 3870
 Phe Pro Ala Ala Gly Arg Ala Leu Ala Ala Leu Ser Val Arg Pro Phe
 3875 3880 3885
 Ala Leu Arg Arg Leu Ser Ala Gly Leu Ser Leu Pro Leu Leu Thr Ser
 3890 3895 3900
 Val Cys Leu Leu Leu Phe Ala Val His Phe Ala Val Ala Glu Ala Arg
 3905 3910 3915 3920
 Thr Trp His Arg Glu Gly Arg Trp Arg Val Leu Arg Leu Gly Ala Trp
 3925 3930 3935
 Ala Arg Trp Leu Leu Val Ala Leu Thr Ala Ala Thr Ala Leu Val Arg
 3940 3945 3950
 Leu Ala Gln Leu Gly Ala Ala Asp Arg Gln Trp Thr Arg Phe Val Arg
 3955 3960 3965
 Gly Arg Pro Arg Arg Phe Thr Ser Phe Asp Gln Val Ala His Val Ser
 3970 3975 3980
 Ser Ala Ala Arg Gly Leu Ala Ala Ser Leu Leu Phe Leu Leu Leu Val
 3985 3990 3995 4000
 Lys Ala Ala Gln His Val Arg Phe Val Arg Gln Trp Ser Val Phe Gly
 4005 4010 4015
 Lys Thr Leu Cys Arg Ala Leu Pro Glu Leu Leu Gly Val Thr Leu Gly
 4020 4025 4030
 Leu Val Val Leu Gly Val Ala Tyr Ala Gln Leu Ala Ile Leu Leu Val
 4035 4040 4045
 Ser Ser Cys Val Asp Ser Leu Trp Ser Val Ala Gln Ala Leu Leu Val

4050 4055 4060
 Leu Cys Pro Gly Thr Gly Leu Ser Thr Leu Cys Pro Ala Glu Ser Trp
 055 4070 4075 4080
 His Leu Ser Pro Leu Leu Cys Val Gly Leu Trp Ala Leu Arg Leu Trp
 4085 4090 4095
 Gly Ala Leu Arg Leu Gly Ala Val Ile Leu Arg Trp Arg Tyr His Ala
 4100 4105 4110
 Leu Arg Gly Glu Leu Tyr Arg Pro Ala Trp Glu Pro Gln Asp Tyr Glu
 4115 4120 4125
 Met Val Glu Leu Phe Leu Arg Arg Leu Arg Leu Trp Met Gly Leu Ser
 4130 4135 4140
 Lys Val Lys Glu Phe Arg His Lys Val Arg Phe Glu Gly Met Glu Pro
 145 4150 4155 4160
 Leu Pro Ser Arg Ser Ser Arg Gly Ser Lys Val Ser Pro Asp Val Pro
 4165 4170 4175
 Pro Pro Ser Ala Gly Ser Asp Ala Ser His Pro Ser Thr Ser Ser Ser
 4180 4185 4190
 Gln Leu Asp Gly Leu Ser Val Ser Leu Gly Arg Leu Gly Thr Arg Cys
 4195 4200 4205
 Glu Pro Glu Pro Ser Arg Leu Gln Ala Val Phe Glu Ala Leu Leu Thr
 4210 4215 4220
 Gln Phe Asp Arg Leu Asn Gln Ala Thr Glu Asp Val Tyr Gln Leu Glu
 225 4230 4235 4240
 Gln Gln Leu His Ser Leu Gln Gly Arg Arg Ser Ser Arg Ala Pro Ala
 4245 4250 4255
 Gly Ser Ser Arg Gly Pro Ser Pro Gly Leu Arg Pro Ala Leu Pro Ser
 4260 4265 4270
 Arg Leu Ala Arg Ala Ser Arg Gly Val Asp Leu Ala Thr Gly Pro Ser
 4275 4280 4285
 Arg Thr Pro Leu Arg Ala Lys Asn Lys Val His Pro Ser Ser Thr
 4290 4295 4300

<210> 3
 <211> 12685
 <212> DNA
 <213> C. Elegans lov-1 gene

<400> 3
 tcaatcttttccacatcgt ttagccgcca cttctggaat ctctttgggtc cagtttcgtg 60
 aatagcagag acaggatcat aggagagtgt gtagttgatg actgtttggt tttggtattg 120
 accttgagtt tggagcattc tgggtggcagc atgatgaagc agattgactt tggcaacagc 180
 gctgtggaat agacggaagt ctttttgagt gtcagcaatt gaaactggag caaaatcttt 240
 tggttcaaga agacccaagc gacgttttgt ctgaaattaa ataacagaaa ttaaagaaca 300
 tctaatagtg agcttgaaaa ataaatacct tgtattttat gtgatcgatt atttcgtaat 360
 cattgggtctg cttctcactg tcattacgaa tttcctcgaa ctcgaacata attatagtga 420

cgttaaagttg caggacgagc ttctgatccgg caatcatata aagcatgata acaacaaaacg 480
 caaattgaga aatcgggttg atagaggtaa catcaagttt tccaagcatt ccaggccaatg 540
 ctgtttgaaa ggttagccatt aagctccgat atctggaacc aattttttaa aattgatttc 600
 ttccaattaa gttttcatcc tcaacctccc attttatttc ctaaaaactg gtacaataca 660
 gagttgaatg tcatgctgaa gaacaggaaa gcaattccaa atgacacaat agctccgaga 720
 gggttatcca gtgtagccgc taatactcca attcttctgt tgaatctcaa gattccgaatc 780
 attttacaag aagtgaagaa taaggctccg gcaagacaat aactgaatac aatctcccaa 840
 tctctctgtt cagtcaaaat aatgtacgaa ttccatttgt ttgcattgaa atcttccatt 900
 gctctatttg tggttcgttg ggggatgggt taggctagga ccgatgcacc agcgagagct 960
 ccaactatca agtccatgaa gttccatggc gagaagtttc tataaaatgt ctttttgaaa 1020
 ctgaagttct cattagaacc accccagtcg cagctgatac acaattttga atggattttct 1080
 cgttggtttc attgttgtta tcaacctgtc ccgcccatac aagtagaaca caatctcttt 1140
 tacaatatg agaactgaga aaaagatgta aagcatctca taatacttga ccacagttcc 1200
 atcgttcccc ctgatttga taagtcttac tgattcaacc caactattag gaagataaat 1260
 tcttgacttt ggaatctcca ccaacaactg taccacggaa aagtagttga ttgagcatt 1320
 gtatgcagag aactcaatga tgactgctcg agtatgata ccgatccatc gttccgaatc 1380
 aagtttattg aagagagtga tgattccgc ttgggtacca gacatactga tagtatatcc 1440
 aactctgaa tagctataca gtaggcctga aactgtttca gtggataatt ctccagaagt 1500
 ctgttaggtg tattcatctg aagcatctgt tccattctcg gattccagtt cggcccaacc 1560
 agcttgcatg tacaaagttt tttcttcgtt tctgaaacct ctctattagt tggaaattga 1620
 agatttttac tcaattgctt gtcaactcct ctccacaatc attgatgtat ccttgaaaact 1680
 gcttgaacat cgtacactct gcaactttct ttgtccgaac ctgccgtatc gtacctatcc 1740
 ccatacttct tgaaacttta tcaattcatg aggcctctcat ccgctatgca ggatttccgt 1800
 cgtaccaaga agccaaaaga gcagtggcca gagattcacg agcccaatcc cagaaatcgt 1860
 cagcatgttg gattgacatg aaagtattgt caccgtagtt cttttgattg atgttcaaga 1920
 ttgtgctcat ctgaaaataa taagttcatc taaatctatg tgcattaaag tctacctcca 1980
 actgatacca atatccatgc cggctcttgc aatagtatgt cagcataacc ataataata 2040
 aagaagcaaa gaaacaaagc atatcacgaa tggttataaa taactgttca tctctcattt 2100
 ttcggttttc agtgtctcgg agctttgtaa catcagcaat ttcggttccc agaccttttt 2160
 cgattttccc atagggattt cctgaatttc agtaatgaat tctgatagct tctttttata 2220
 aaacttactc aagaacgtct cagctggctt agctcttagc aatgcttctt ccaacttggt 2280
 aatgatttta tgactctttc tggttttcaa aattaaaaac gcccaaatca atcccttaat 2340
 tggctcgaac accactgccc atagaatcag actgatcaga aatcggatat agaaagagtt 2400

gggtaaatca tccatcaagc tcattccagc tccagaaata taaataagac ccattgagaac 2460
 tggaaataact atgatggtaa gtgccatccc agccatgaac atgggcaatg aaccactatt 2520
 atccttgaat tccggatcct ctctcttttg ttttttgtag tagtaatgtt cactgtggga 2580
 aagacatttg gtgcataata aaatgtgcaa tgagttgagg aaagtgataa gaacaccgaa 2640
 tccaaactccg aatgcaatat cttttatagt gaaagtgaac tggagagac ctttcgaatc 2700
 actgataatc gaattatccg ttttcagaat tgtgatgcta atcatgctga ccacaacaag 2760
 tgagaagatg atactgacag aatagtcctg ctttgacact cgatccctca accgattgoc 2820
 tccaccagta aacatggcaa accaggaaat tgtttgagoc agcatatgca taactcattga 2880
 ctcacccaaa aaccttccgt tatactccac tggcgctagt ctttcagctt ctccgtctcc 2940
 gtttttagtt ccaagccaat tgttgaaaag gaagtagtag atatcctgag tctgtagatc 3000
 tttcacatt attcgattgc aataccacga ctctcggtag tctagaccag catcgtaag 3060
 ccagagtctc atgtattcca actcgccaag agggctgaaa tattaaattt ggtaaatgat 3120
 ttttgatttg aaaacttgaa ttagtccatc aaaaacccaa accaagttagg gggataaaaa 3180
 aaactacacg tccaatctat aattagctca actcacactt gaaccaatca ggttgctccg 3240
 aatttgcata attggttgaa acgtgtgtgg agctaactgg tattatttat ttttaattatt 3300
 tttttatttg taaaaatcag cgtcttctaa cttacaatgc cgttgctcat acaaatcgat 3360
 cagtgggttc ccattgaaaac ggaaactccc aattaccatc ttctctgat ctgaacgatc 3420
 ggaaaatctg atccccctca tttccagata aattgaaaac tatcgtaacta tccgtagttg 3480
 caaacattcg atatccagtc tccacggcaa tccatataat gtatccatca tgaggctcat 3540
 tgtctttcag aaaacgaagt ctcccgctg atgcactctt acgttgacag atgattgcac 3600
 tgatggtaag acatccgtaa actactagca tgaaaacagc ggcaatcatc actttcacat 3660
 tcttttcgat ttcattcaca ttataattgt aagagaaatc tgcataata gttggattga 3720
 atgcaccaac agagaacatt gttaaatgat cagttgaaca attaacgaac tgcattcctt 3780
 gtccatcact tggatacatt ctttcagaat tgaagacatc cgatgttttc tgatagaagt 3840
 aacatccttt actcaatgca gcgacttgat aatccattgg tacacttcgt gcaaatgacc 3900
 actgcattga atcgtaactga ccatagttta caatatccga actatttcca gtgttagtag 3960
 agctattttc ttttccaaat ccaataaaga aaagtccagt gttgttgatc aaatttccgg 4020
 cggtgacaaa ataattgctt gtcttgttca atgtgttcaa gtcaaagatc cattcatgat 4080
 ttgattcaag tgggcccagga agactttgga atgatgagaa catgtaggtg tcatcgttgt 4140
 ttggaatttc atagtcttga gatgcaataa tctctacttg aagcgagttg ttccagttag 4200
 tggttcgaaa agcatgaaga tctaatactt gataactggc aaagtctcct tgttgcatca 4260
 aagttagcac tgcacatcc ccaactcccc tgcggttcac ataaattgga gcagtagttc 4320
 cggttattct aaaaacattt taacttatat tggaaaaatt ataggttatt caataaactt 4380
 acggaatgat tatctgatcc tcatctttga tatgtgcttc aagtgcacct gaggtaatta 4440

acatatcaaa gttatccaca taagttctcg ggtttgtggc atagcaaac aatccaaact 4500
 gaatcagtggt tttatctgtg atctcagcag tattcagagt tgaagcgggc gatggaagtt 4550
 tgaatgccc ttcttcacag ttttgagttt ttctacaaat atttgatgca tcatcgataa 4620
 cgattacat ttccggttcg tcgacgctat tctaaaaatt tgattgacat tagtgtaaac 4630
 tghtaactttt tgattacoga gtagtcataa ggtaagttgc cagttgctat agctctagct 4740
 gctagcgtgt tttccagggt atctagtgtg gatgcaagtt gggtggctag atttctggcc 4800
 tgaatagaat atgaataatt ccaaaactca aaagttttca aaactcaga tgttctctcg 4850
 gaacattttt gtgacgtaag cagccattc ttctgaagtc atttctcca cgtacacaat 4920
 attgtctgga tcaactggta gcacgttgta caaggaatca tagttatctg ttgcataatt 4930
 caaattggca gctagatcag aagaaagagg attgtcaga gcaatctcca acgctgatgt 5040
 taaagatccc gcaattgaaa ggagagagtt ggatgtctga aaattattat tatgacatct 5100
 accaaagttt agtgtatgaa tacatcatta atctctgcat ctgtcatagt cattccatta 5150
 tttgtaagaa aatcttgagt attgctaaga atcgtttcaa tttgttctg ggattcttga 5220
 ttcatcaaac ttgcggcac aatatctgga tatttaaaat gatttgtatg catttctgta 5230
 tttatttttt tgagttacca attttagttc ccgaaacatc gccttcctgc gaaacttctc 5340
 cattctgaat tacatatcca ttcgttccat ttctaataat caacctcca tcatctcca 5400
 tttctctgtg gtcagagaat tccattaaag ttctcaaat taaagtctct atacaaaata 5450
 tccattcgag actatactta caccgtatga gtttgagaac cagatgtgta agttccgggt 5520
 gttatctgaa taccgtaacc tccaagcgc aaagaaacaa tattgtaggt gggagaactg 5530
 atcaccagag ttgtccaga gtttgacaaa taaagggcgg tggagtctac gaaaaatgag 5640
 tagccgctga catcgatgga tgacgctttg gagaagaata gctgcaagaa aagttatttt 5700
 gatattaaca actcatcagc aaaagtctta cagttccga aacggtgaca agtgtttgtt 5750
 ccttaatcaa tgattgcaaa tcatctttgg tgtacgaagc cgtcggtgaa agagtagata 5820
 ttgagacagt ttgagataaa gcagagacag ctggaattcc aatgtctctc ggagacagca 5830
 ccattcgcat tgaataactt ccgcggtttg agaacacaag tggagaagta gtttgattga 5940
 ataaagcaag gttgacgtct gaaattttta gctcataac cagaccactc ccattgcata 6000
 ctactctca ataactatcg gcactctaat caactctta accgtgctcg ctgaggatcc 6050
 atctgatgcc acaaccgtga agctgtcagc attgtatgag aaaatggcat tatctgtgtc 6120
 tacttcaaca aactttctcg ttccattcgt gcaagttatt gtgtaagtaa ctccaccata 6180
 tgatgcagtt actactagtc ctctagatac aatattttga gtagaatgca gttttatcat 6240
 tgttcagat tcaatctgaa taaaaattga aaaatttcat gtgctctacg atttataaat 6300
 ttaccaataa tatgtattga acttctgaga ctgggatatt gaacgaaaat gaagtgttct 6350
 tctgacttgt ggatctact gttgaattgt cataagtgac attccaagca gttacataca 6420

tatcattata actatattcc cctgattgaa catctccaac tccactcatt gtagccatta 6430
 tatttcacaga aaccagtgat cgtttgtcat ctattttato ctgctcaact tgcacagtc 6510
 ttccatttgt tgcgaacatt ccttgggatto cgatggcagg tgcctagcata gtgtccgct 6590
 gaacagtttc acataaaacta caatgttcta tactcaaaaa gtcttacagt aacggtatct 6670
 ccactaagct gtttataaga tgcctgggta tctctgttaa ggtatattgt agatccgtaa 6750
 atgctgagct gtaagttgaa aacaattaac tctcccaacc atcattttct taccgtacac 6830
 ctaggcgata ccaatgtata tcccgatgca ataacatato caaaaataac cggctaaagt 6910
 ccactttttg attttctaact tgatactccc aaggcataga ctgtacttcc ttgagagaca 7000
 agatccagag aagacagtac ggagtcacag gattgtgogg aagtcatact gacatttct 7080
 agtttagtga taacttttgc catttcgtcg gccaatccg aatttgttgt tgcattattg 7160
 tcttgcaatg ttttcaaaac atcaaacctt gacatatttc cgactccagg gattttgagg 7240
 gtatttgaga gcaaaccttg agcaacttcc actagatctg cggcaggtag agatgagatt 7320
 tgattgagaa gagagctgct tgtgtttaga gagttgttgg atgcagatcc atccattatc 7400
 ccagccagtt ggttcattac atcagctttt tgagcatcta tgatcgcttg tccagctgca 7480
 gaaattggag aaactgttgc aagagagctg cgagttttag tagttcttct agatggttgc 7560
 ggggtagcag agaatgcacc atttgcacca gaggaatcgg aagatccaga cgtatccgag 7640
 ccagatgagc tcttggttga aacaccagaa gatccatttg aatctgaccc tgagccagat 7720
 gtactaccac cgtctcctaa atgggatccct ggtgtggtag ctgttccaga tctgtacca 7800
 tctccattca atgccttctg ttttcacag tttaccctgt ccgaaccctg cctgaagac 7880
 cccctgacc cacttccaga agcgtctgtt cctgatccac cagcccccga tcccttgat 7960
 gattgtccac ttcagatcc agaagtgtgt gatctgactg catcaccctg gctaagagtt 8040
 gtccagatc ctctgaacc tgttccacca gttcccccag tagctccagt tccaccggtt 8120
 ttgcccagg catcatcca cgagctgaca gtggttgggt gggtttctaa aaattgaatt 8200
 ttatgaaaaa aaaacagtaa tgcgcttacc agtagttgta gttggaagaa ctacattcat 8280
 agtaaagatg tgaactggcag attctccaga tgcacgattg gtaacattaa ttaagaattc 8360
 ataagttcca gttgcaggta caaagctggt cattgggttg aaagatacgg aggcactata 8440
 tgcaccattt tttccaactg taatatagta cataaatgaa aattgataac gttgctaact 8520
 agggagtgt ctttgtactc catggaaata tgagtcggaa aactactttt cgggtagttt 8600
 atgtcatttt ctacacgatt ctgaaagaaa atcctgccgg ttttgggttt tagtgtgaaa 8680
 agtttgcgtt tgaaaatacc accctaaatt cagttgattt aacactacgc gacccatatt 8760
 tcatgtgcaa cgggaaagcc aagtacactg aaaactcact ttcaaatttt caaagcaaag 8840
 tcataatttc ggtggtccag tggataacgg cgggagcggc gccagttttc agtgtacttg 8920
 gctttccctg tgcacatgaa atatgggtcg cgtagtgtta aatcaactga atttagggtg 9000
 gtattttcaa acgcaaactt ttcacactaa aacccaaaac cggcaggatt ttctttcaga 9080

atcgtgtaga aaatgatatc aactacogga aaagtagttt tccgactcat atttccatgg 8520
 agtacaaaaa gcactcccta gtaggcaaaa tctcacactc tgtcaagcaa ttgctttcct 8580
 tcagtaaaaac aaatggctga gaagaatcgt ttcgacattg acaggtcata gcagtcttaa 8640
 aatattaagg ttttttttaa gtaagattga tttgaatata ttacogttaa agtcoggggc 8700
 tctttgtttg gaattggtgt aatataaaat ggattttcag agtaatacac tgtctcgttc 8760
 catgtcaaat tttccaatac aaagtogtaa ggggtttctg taottgogtc tggatccaca 8820
 gttgttgttc tcatgctacc agatgcactt gatgaatcgg atggtgtttg agataaacca 8880
 gatgaatctg aagtggatgg agtcgagtta gatgagtcta tgggtggtgga atctgaagtt 8940
 gtgcacagaat ccgatgttga tgtagaactg tcttgtgaag catcagtcgt gctagcttcc 9000
 aaagtogaag tagattcaat tctgaagtt gtggaaatgg ctgaactttc ggaagaagat 9060
 ccagttgaag ttgtactggt aacttcggat gtacttggtg aatccgcaac aacattcaat 9120
 gtgaaaatgt ggttgaccac ttgcattgtc gtcaaatccg tcatgttgat tcgaaactca 9180
 taggtgcaca tgccaacaag aaatgtttca attggttgaa ttggaatatt agaagaataa 9240
 gttgogtcta ggacogtgtt tcttggaat tgcgtttaat tcaataattt caaaaagttt 9300
 gttaatccta cagtagttta agcaagtga tctcttaatt ataagaaaag gctcagtaga 9360
 cacattctcg cattcaaatg tttggcttct ctaaaatcat tegtttcatt tggctcaacg 9420
 atttattaaa ccgcctcag taggagtaat ggcattagtt ggtaaaggta caatgttgat 9480
 gctgtcttca ttgtgacgtg tctcgttcca tgacaatcca ctgtccaaga tgaaatcgaa 9540
 ctgatogaat gacaaagtgg aagttgaatc agcagtggtg gaactcggac tggagaagat 9600
 tgtggatgtg tccgagatgg tagttgtact ggaatcagta gtactttctt cggaggttgt 9660
 tgtggattct tgtaaagtag ttgtgctccc agccgacgtt gtctgagaat cgtttgaacg 9720
 tgtggatgac ggtctctgta ctgtggatgt tgataatgtg gaagatggag ttgatggtgt 9780
 agatgacgta gagcttgcta cagcagatgt agaagattcc gactctatgg tggtggaaga 9840
 atattcctga atataaacgt tcgcatacgt gtagtaaaact tttttatcgt cggttgtcat 9900
 agttgctctg aaggtgtagt ttccaggacc aacgaatgtg ctagcagggg acgtccctcc 9960
 gagacgtggc attgatacac tttctgaata tattgaaaaa atatgtgtaa aaaatctaaa 10020
 taactatcgc ccgaaaaagg tttgcttttt ttccgatttg aagtttttat agaaacgttt 10080
 tcagaattaa agattttgcc tgtctctaatt ttataataag tctttataaa caattactcg 10140
 tgaaacatgc tccgtcttta gtggttgaaa cataattaga agatgtaggg cttgtacatt 10200
 cgatggatgt ttgatactga aatacagtgt tacatttgaa taatgcagtc ttcaatattg 10260
 tacttactcc aatgattccg aggcoggaat taagcgttag gttcacactt gtggaatcat 10320
 agaacgttgt agttgccttt tctacaaaat agaagtcctg attagttccg tccgagctag 10380
 tagttgtctc agattttggt gttgttgaaac tttgctgagt agacgtggat gattgagtag 10440

aagaagcggg gcttgacaca gacgaagaag ccgttgaact tggagtggaa gaagaactcg 10500
atggcccagt agttgatgtt gaaggagcag ttgtgctggt tgttactgta gaogaaggag 10600
atgtcgatgt ggactcagt cttgttgggg ttgttacagt agtcgaggag cttgaactag 10700
atgtacccgt agaagtcaca ggggatgttg acggggaagt agttacagta ctgtcgatg 10800
gttcgggtgt tgaagttgaa gcagtcgagg tggtaaaagt agtggttggt tcagttgttg 10900
taacagtaga agatgttagat gtaactttctg ttgtgggtgt ggaagtagat ggttcttcgg 11000
tagtagtgga tgtcaacata gtggtggtga aagtggtaga cgtagtggtc tegtcaaggt 11100
aggagcaaat cgcattatcc gggagagacg acaaagttgc tgaatatctt cgttaaggat 11200
tttctggata actaacaatg cacaacaagg tgatcggtaa tagtgaactgc tttgttttac 11300
cctgagcaaa ctgtaattgt ataaaaatctg aaatattggc aatacaaacg ggtttgaaga 11400
aaattattaa caattttatt cctgcctctc aatcataaca gcaaatctct gtttgcttgc 11500
aattattatt gtggttcga aactcacatg tgatttcggg tgttgtagtg gttggaatac 11600
ttgtaactcag tgtggtggag ctcggcgagc ttgtgattgt gctagaactc tegtgagttg 11700
tgctagttag tgatgtcgac gtggatgctg tggaaagtga cgttggttag gtggattoga 11800
tggtggtgga tgttgatgga gttgatgtgc ttgtgctcat tgcggtagtc accgtaactg 11900
tacttgttgg gaaggttggt gtagatgtca cggtaactag cagggttgtg gtagttggag 12000
ttgtagtoga ggttgaagtt gactcaatag tgtagtoga agtattatca ccttggataa 12100
aaatgaaagt aaacactatc tgagaaatcg tactcacagc gtctcgtttc attcttctca 12200
aagtaggtga tccagaagtc ctcattgctaa actgttttgc cctgacaccc ttaagtacct 12300
gcccataata ataacactct tgactatcac tgatagctgt gctcttttca gaacgatctg 12400
aaatactggt tagccaatgt tcattgagcaa ttaagaactg acaaggctgc ttgcacattc 12500
ttctgcata ctcttcgttg atctctccgc tctcacactt ctctcggtag cccagcaacg 12600
ccatctcggt tccaccgact tggagccacc atagggagcc atcacatctg tcgataccgt 12700
catctgagaa agagtttcta ttaaaatggt agaaacacat agcactacat atgcaaataa 12800
cgtttcacca gattcagaat gcgcaattca tgctatctc atagcctacc tatgtgtcta 12900
cctgagtatc tacttgagta ccttgcaaag aagattaatc ggcacaaacc aagtcaagac 13000
tttgttggca taggtcttcc aggtgagtaa cgcgacatt atacataggt accacaaaaa 13100
ccttccccc aaataaatcc ttaccataac aaacttcata ttctgcctcc acagcaatac 13200
tgatctcatc gtcacattc acttcattca aagtaatcca agttgagttc aaaaagagtc 13300
cgacaagcct ggtctctggt tggatgcagt tgtgaatctg aataggaaca acaaggtttt 13400
acaactaaaa aaatacacga ctaaccaatt ccaaacttga aacttcgta accttgttct 13500
caactgaaag tctattcaat ccgcagctca atttgatttt aacgactcct tgtgaattcc 13600
ttggaactcc tccaattggt gtgtcatcgt tgtctaactg aaaagttgcg atcccgtaaa 13700
gaagttggta atgcaatcca tcaatttgta tcttaaaagt agttttatcc agcttttctc 13800

totgagattt ttcaatcacc gccgatattg ccagtagcaa tagaacaaaag aagtttgact 12540
 ttttcatcda atgagctgga aggttatatt gtagaagttt tgtaaaaatt cgcctgaaaa 12600
 caaaaatgaa tttagagcag aaaagacaa aactgaaaaa tgaagttgtc gaaaagcgaa 12660
 aagggggggt gaatgaagg accat 12685

<210> 4
 <211> 3173
 <212> PRT
 <213> C. Elegans Ldv-1 protein

<400> 4
 Met Val Leu Arg Phe Ser Pro Pro Phe Arg Phe Ser Thr Thr Ser Phe
 1 5 10 15
 Phe Ser Cys Cys Leu Phe Cys Ser Glu Phe Ile Phe Val Phe Arg Arg
 20 25 30
 Ile Phe Thr Lys Leu Leu Gln Asp Asn Leu Pro Ala His Trp Met Lys
 35 40 45
 Lys Ser Asn Phe Phe Val Leu Leu Leu Leu Ala Ile Ser Ala Ile Gln
 50 55 60
 Ile Asp Gly Leu His Tyr Gln Leu Leu Asp Gly Ile Ala Thr Phe Arg
 65 70 75 80
 Leu Asp Asn Asp Asp Thr Thr Ile Gly Gly Val Pro Arg Asn Ser Gln
 85 90 95
 Gly Val Val Lys Ile Lys Leu Ser Cys Gly Leu Asn Arg Leu Ser Val
 100 105 110
 Glu Asn Lys Val Thr Glu Val Ser Ser Leu Glu Leu Ile His Asn Cys
 115 120 125
 Ile Gln Thr Glu Thr Arg Leu Val Gly Leu Phe Leu Asn Ser Thr Trp
 130 135 140
 Ile Thr Leu Asn Glu Val Asn Asp Asp Asp Glu Ile Ser Ile Ala Val
 145 150 155 160
 Glu Ala Lys Tyr Glu Val Cys Tyr Asp Asp Gly Ile Asp Arg Cys Asp
 165 170 175
 Gly Ser Leu Trp Trp Leu Gln Val Gly Gly Asn Glu Met Ala Leu Leu
 180 185 190
 Gly Tyr Arg Glu Lys Cys Glu Ser Gly Glu Ile Asn Glu Glu Tyr Ala
 195 200 205
 Arg Arg Met Cys Lys Arg Pro Tyr Arg Ser Glu Lys Ser Thr Ala Ile
 210 215 220
 Ser Asp Ser Gln Gly Val Tyr Tyr Asp Gly Gln Val Leu Lys Gly Val
 225 230 235 240
 Arg Ala Lys Gln Phe Ser Met Arg Thr Ser Gly Ser Pro Thr Leu Arg

245										250					255				
Arg	Met	Lys	Arg	Asp	Ala	Gly	Asp	Asn	Thr	Cys	Asp	Tyr	Thr	Ile	Glu				
			260					265					270						
Ser	Thr	Ser	Thr	Ser	Thr	Thr	Thr	Pro	Thr	Thr	Thr	Thr	Thr	Val	Thr	Ser			
		275					280						285						
Thr	Val	Thr	Ser	Thr	Thr	Thr	Val	Pro	Thr	Ser	Thr	Ser	Thr	Val	Thr				
	290					295						300							
Thr	Ala	Met	Ser	Thr	Ser	Thr	Ser	Thr	Pro	Ser	Thr	Ser	Thr	Thr	Ile				
305					310					315					320				
Glu	Ser	Thr	Ser	Thr	Thr	Phe	Thr	Ser	Thr	Ala	Ser	Thr	Ser	Thr	Ser				
			325					330						335					
Ser	Thr	Ser	Thr	Thr	Gln	Gln	Ser	Ser	Ser	Thr	Ile	Thr	Ser	Ser	Pro				
			340				345						350						
Ser	Ser	Thr	Thr	Leu	Ser	Thr	Ser	Ile	Pro	Thr	Thr	Thr	Thr	Pro	Glu				
		355					360					365							
Ile	Thr	Ser	Thr	Leu	Ser	Ser	Leu	Pro	Asp	Asn	Ala	Ile	Cys	Ser	Tyr				
	370					375				380									
Leu	Asp	Glu	Thr	Thr	Thr	Ser	Thr	Thr	Phe	Thr	Thr	Thr	Met	Leu	Thr				
385					390				395					400					
Ser	Thr	Thr	Thr	Glu	Glu	Pro	Ser	Thr	Ser	Thr	Thr	Thr	Thr	Glu	Val				
				405					410					415					
Thr	Ser	Thr	Ser	Ser	Thr	Val	Thr	Thr	Thr	Glu	Pro	Thr	Thr	Thr	Leu				
			420				425						430						
Thr	Thr	Ser	Thr	Ala	Ser	Thr	Ser	Thr	Thr	Glu	Pro	Ser	Thr	Ser	Thr				
		435					440					445							
Val	Thr	Thr	Ser	Pro	Ser	Thr	Ser	Pro	Val	Thr	Ser	Thr	Val	Thr	Ser				
	450					455				460									
Ser	Ser	Ser	Ser	Ser	Thr	Thr	Val	Thr	Thr	Pro	Thr	Ser	Thr	Glu	Ser				
465					470					475				480					
Thr	Ser	Thr	Ser	Pro	Ser	Ser	Thr	Val	Thr	Thr	Ser	Thr	Thr	Ala	Pro				
			485					490						495					
Ser	Thr	Ser	Thr	Thr	Gly	Pro	Ser	Ser	Ser	Ser	Ser	Thr	Pro	Ser	Ser				
		500					505						510						
Thr	Ala	Ser	Ser	Ser	Val	Ser	Ser	Thr	Ala	Ser	Ser	Thr	Gln	Ser	Ser				
	515						520						525						
Thr	Ser	Thr	Gln	Gln	Ser	Ser	Thr	Thr	Thr	Lys	Ser	Glu	Thr	Thr	Thr				
	530				535					540									
Ser	Ser	Asp	Gly	Thr	Asn	Pro	Asp	Phe	Tyr	Phe	Val	Glu	Lys	Ala	Thr				
545					550					555				560					
Thr	Thr	Phe	Tyr	Asp	Ser	Thr	Ser	Val	Asn	Leu	Thr	Leu	Asn	Ser	Gly				
			565						570				575						
Leu	Gly	Ile	Ile	Gly	Tyr	Gln	Thr	Ser	Ile	Glu	Cys	Thr	Ser	Pro	Thr				
		580					585						590						
Ser	Ser	Asn	Tyr	Val	Ser	Thr	Thr	Lys	Asp	Gly	Ala	Cys	Phe	Thr	Lys				

595					600					605					
Ser	Val	Ser	Met	Pro	Arg	Leu	Gly	Gly	Thr	Tyr	Pro	Ala	Ser	Thr	Phe
610						615					620				
Val	Gly	Pro	Gly	Asn	Tyr	Thr	Phe	Arg	Ala	Thr	Met	Thr	Thr	Asp	Asp
625					630					635					640
Lys	Lys	Val	Tyr	Tyr	Thr	Tyr	Ala	Asn	Val	Tyr	Ile	Gln	Glu	Tyr	Ser
					645					650					655
Ser	Thr	Thr	Ile	Glu	Ser	Glu	Ser	Ser	Thr	Ser	Ala	Val	Ala	Ser	Ser
					660					665					670
Thr	Ser	Ser	Thr	Pro	Ser	Thr	Pro	Ser	Ser	Thr	Leu	Ser	Thr	Ser	Thr
					675					680					685
Val	Thr	Glu	Pro	Ser	Ser	Thr	Arg	Ser	Ser	Asp	Ser	Thr	Thr	Thr	Ser
					690					695					700
Ala	Gly	Ser	Thr	Thr	Thr	Leu	Gln	Glu	Ser	Thr	Thr	Thr	Ser	Glu	Glu
705					710					715					720
Ser	Thr	Thr	Asp	Ser	Ser	Thr	Thr	Thr	Ile	Ser	Asp	Thr	Ser	Thr	Ser
					725					730					735
Thr	Ser	Ser	Pro	Ser	Ser	Thr	Thr	Ala	Asp	Ser	Thr	Ser	Thr	Leu	Ser
					740					745					750
Val	Asp	Gln	Phe	Asp	Phe	Ile	Leu	Asp	Ser	Gly	Leu	Ser	Trp	Asn	Glu
					755					760					765
Thr	Arg	His	Asn	Glu	Asp	Ser	Ile	Asn	Ile	Val	Pro	Leu	Pro	Thr	Asn
					770					775					780
Ala	Ile	Thr	Pro	Thr	Glu	Arg	Ser	Gln	Thr	Phe	Glu	Cys	Arg	Asn	Val
785					790					795					800
Ser	Thr	Glu	Pro	Phe	Leu	Ile	Ile	Lys	Glu	Ser	Thr	Cys	Leu	Asn	Tyr
					805					810					815
Ser	Asn	Thr	Val	Leu	Asn	Ala	Thr	Tyr	Ser	Ser	Asn	Ile	Pro	Ile	Gln
					820					825					830
Pro	Ile	Glu	Thr	Phe	Leu	Val	Gly	Ile	Gly	Thr	Tyr	Glu	Phe	Arg	Ile
					835					840					845
Asn	Met	Thr	Asp	Leu	Thr	Thr	Met	Gln	Val	Val	Ser	His	Ile	Phe	Thr
					850					855					860
Leu	Asn	Val	Val	Ala	Asp	Ser	Thr	Ser	Thr	Ser	Glu	Val	Thr	Ser	Thr
865					870					875					880
Thr	Ser	Thr	Gly	Ser	Ser	Ser	Glu	Ser	Ser	Ala	Ile	Ser	Thr	Thr	Ser
					885					890					895
Gly	Ile	Glu	Ser	Thr	Ser	Thr	Leu	Glu	Ala	Ser	Thr	Thr	Asp	Ala	Ser
					900					905					910
Gln	Asp	Ser	Ser	Thr	Ser	Thr	Ser	Asp	Ser	Gly	Thr	Thr	Ser	Asp	Ser
					915					920					925
Thr	Thr	Ile	Asp	Ser	Ser	Asn	Ser	Thr	Pro	Ser	Thr	Ser	Asp	Ser	Ser
					930					935					940
Gly	Leu	Ser	Gln	Thr	Pro	Ser	Asp	Ser	Ser	Ser	Ala	Ser	Asp	Ser	Met

945	950	955	960
Arg Thr Thr Thr Val Asp Pro Asp Ala Ser Thr Glu Thr Pro Tyr Asp			
	965	970	975
Phe Val Leu Glu Asn Leu Thr Trp Asn Glu Thr Val Tyr Tyr Ser Glu			
	980	985	990
Asn Pro Phe Tyr Ile Thr Pro Ile Pro Asn Lys Glu Pro Gly Ala Leu			
	995	1000	1005
Thr Thr Ala Met Thr Cys Gln Cys Arg Asn Asp Ser Ser Gln Pro Phe			
	1010	1015	1020
Val Leu Leu Lys Glu Ser Asn Cys Leu Thr Glu Phe Gly Lys Asn Gly			
1025	1030	1035	1040
Ala Tyr Ser Ala Ser Val Ser Phe Asn Pro Met Thr Ser Phe Val Pro			
	1045	1050	1055
Ala Thr Gly Thr Tyr Glu Phe Leu Ile Asn Val Thr Asn Arg Ala Ser			
	1060	1065	1070
Gly Glu Ser Ala Ser His Ile Phe Thr Met Asn Val Val Leu Pro Thr			
	1075	1080	1085
Thr Thr Thr Glu Thr Pro Pro Thr Thr Val Ser Ser Ser Asp Asp Ala			
	1090	1095	1100
Gly Gly Lys Thr Gly Gly Thr Gly Ala Thr Gly Gly Thr Gly Gly Thr			
1105	1110	1115	1120
Gly Ser Gly Gly Ser Ala Thr Thr Leu Ser Thr Gly Asp Ala Val Arg			
	1125	1130	1135
Ser Thr Thr Ser Gly Ser Gly Ser Gly Gln Ser Ser Thr Gly Ser Gly			
	1140	1145	1150
Ala Gly Gly Ser Gly Thr Thr Ala Ser Gly Ser Gly Ser Gly Gly Ser			
	1155	1160	1165
Ser Gly Thr Gly Ser Asp Gly Val Asn Ser Gly Lys Thr Thr Ala Leu			
	1170	1175	1180
Asn Gly Asp Gly Thr Gly Ser Gly Thr Ala Thr Thr Pro Gly Ser His			
1185	1190	1195	1200
Leu Gly Asp Gly Gly Ser Thr Ser Gly Ser Gly Ser Asp Ser Asn Gly			
	1205	1210	1215
Ser Ser Gly Val Ser Thr Lys Ser Ser Ser Gly Ser Asp Thr Ser Gly			
	1220	1225	1230
Ser Ser Asp Ser Ser Gly Ala Asn Gly Ala Phe Ser Ala Thr Ala Gln			
	1235	1240	1245
Pro Ser Thr Arg Thr Thr Lys Thr Arg Ser Ser Leu Ala Thr Val Ser			
	1250	1255	1260
Pro Ile Ser Ala Ala Glu Gln Ala Ile Ile Asp Ala Gln Lys Ala Asp			
1265	1270	1275	1280
Val Met Asn Gln Leu Ala Gly Ile Met Asp Gly Ser Ala Ser Asn Asn			
	1285	1290	1295
Ser Leu Asn Thr Ser Ser Ser Leu Leu Asn Gln Ile Ser Ser Leu Pro			

1300	1305	1310
Ala Ala Asp Leu Val Glu Val Ala Gln Ser Leu Leu Ser Asn Thr Leu 1315 1320 1325		
Lys Ile Pro Gly Val Gly Asn Met Ser Ser Val Asp Val Leu Lys Thr 1330 1335 1340		
Leu Gln Asp Asn Ile Ala Thr Thr Asn Ser Glu Leu Ala Asp Glu Met 1345 1350 1355 1360		
Ala Lys Val Ile Thr Lys Leu Ala Asn Val Asn Met Thr Ser Ala Gln 1365 1370 1375		
Ser Leu Asn Ser Val Leu Ser Ser Leu Asp Leu Ala Leu Lys Gly Ser 1380 1385 1390		
Thr Val Tyr Thr Leu Gly Val Ser Ser Thr Lys Ser Lys Asp Gly Thr 1395 1400 1405		
Tyr Ala Val Ile Phe Gly Tyr Val Ile Ala Ser Gly Tyr Thr Leu Val 1410 1415 1420		
Ser Pro Arg Cys Thr Leu Ser Ile Tyr Gly Ser Thr Ile Tyr Leu Thr 1425 1430 1435 1440		
Gly Asp Thr Arg Ala Ser Tyr Lys Gln Leu Asp Gly Asp Thr Val Thr 1445 1450 1455		
Ala Asp Thr Met Leu Ala Ala Ala Ile Gly Ile Gln Gly Met Phe Ala 1460 1465 1470		
Thr Asn Gly Arg Thr Val Gln Val Glu Gln Asp Lys Ile Asp Asp Lys 1475 1480 1485		
Arg Ser Leu Val Ser Gly Asn Ile Met Ala Thr Met Ser Gly Val Gly 1490 1495 1500		
Asp Val Gln Ser Gly Glu Tyr Ser Tyr Asn Asp Met Tyr Val Thr Ala 1505 1510 1515 1520		
Trp Asn Val Thr Tyr Asp Asn Ser Thr Val Gly Ser Thr Ser Gln Lys 1525 1530 1535		
Asn Thr Ser Phe Ser Phe Asn Ile Pro Val Ser Glu Val Gln Tyr Ile 1540 1545 1550		
Leu Leu Ile Glu Ser Gly Thr Met Ile Lys Leu His Ser Thr Gln Asn 1555 1560 1565		
Ile Val Ser Arg Gly Leu Val Val Thr Ala Ser Tyr Gly Gly Val Thr 1570 1575 1580		
Tyr Thr Ile Thr Cys Thr Asn Gly Thr Gly Lys Phe Val Glu Val Asp 1585 1590 1595 1600		
Thr Asp Asn Ala Ile Phe Ser Tyr Asn Ala Asp Ser Phe Thr Val Val 1605 1610 1615		
Ala Ser Asp Gly Ser Ser Ala Ser Thr Val Lys Lys Leu Ile Gln Met 1620 1625 1630		
Pro Ile Val Ile Glu Asn Val Asn Leu Ala Leu Phe Asn Gln Thr Thr 1635 1640 1645		
Ser Pro Leu Val Phe Ser Asn Ala Gly Ser Tyr Ser Met Arg Met Val 1650 1655 1660		

Leu Ser Pro Gln Asp Ile Gly Ile Pro Ala Val Ser Ala Leu Ser Gln
 1665 1670 1675 1680
 Thr Val Ser Ile Ser Thr Leu Ser Pro Thr Ala Ser Tyr Thr Lys Asp
 1635 1690 1695
 Asp Leu Gln Ser Leu Ile Lys Glu Gln Thr Leu Val Thr Val Ser Gly
 1700 1705 1710
 Thr Thr Ser Asn Ser Leu Leu Ser Ile Ala Gly Ser Leu Thr Ser Ala
 1715 1720 1725
 Leu Lys Ile Ala Leu Asp Asn Pro Leu Ser Ser Asp Leu Ala Ala Asn
 1730 1735 1740
 Leu Lys Tyr Ala Thr Asp Asn Tyr Asp Ser Leu Tyr Asn Val Leu Pro
 1745 1750 1755 1760
 Ser Asp Pro Asp Asn Ile Val Tyr Val Glu Glu Met Thr Ser Glu Glu
 1765 1770 1775
 Trp Ala Ala Tyr Val Thr Lys Met Phe Gln Lys Asn Ile Ala Lys Asn
 1780 1785 1790
 Leu Ala Asn Gln Leu Ala Ser Thr Leu Asp Thr Leu Glu Asn Thr Leu
 1795 1800 1805
 Ala Ala Arg Ala Ile Ala Thr Gly Asn Leu Pro Tyr Asp Tyr Ser Asn
 1810 1815 1820
 Ser Val Asp Gly Thr Gly Met Val Ile Val Ile Asp Asp Ala Ser Asn
 1825 1830 1835 1840
 Ile Val Gly Lys Thr Gln Asn Cys Glu Glu Trp Ala Phe Lys Leu Pro
 1845 1850 1855
 Ser Pro Ala Ser Thr Leu Asn Thr Ala Glu Ile Thr Asp Lys Thr Leu
 1860 1865 1870
 Ile Gln Val Gly Leu Val Cys Tyr Ala Thr Asn Pro Arg Thr Tyr Val
 1875 1880 1885
 Asp Asn Phe Asp Met Leu Ile Thr Ser Gly Ala Leu Glu Ala His Ile
 1890 1895 1900
 Lys Asp Glu Asn Gln Ile Ile Ile Pro Ile Thr Gly Thr Thr Ala Pro
 1905 1910 1915 1920
 Ile Tyr Val Asn Gly Arg Gly Ser Glu Asp Asp Ala Val Leu Thr Leu
 1925 1930 1935
 Met Gln Gln Gly Asp Phe Ala Ser Tyr Gln Ile Leu Asp Leu His Ala
 1940 1945 1950
 Phe Arg Thr Thr Asn Trp Asn Asn Ser Leu Gln Val Glu Ile Ile Ala
 1955 1960 1965
 Ser Gln Asp Tyr Glu Ile Pro Asn Asn Asp Asp Thr Tyr Met Phe Ser
 1970 1975 1980
 Ser Phe Gln Ser Leu Pro Gly Pro Leu Glu Ser Asn His Glu Trp Ile
 1985 1990 1995 2000
 Phe Asp Leu Asn Thr Leu Asn Lys Thr Ser Asn Tyr Phe Val Thr Ala
 2005 2010 2015

Gly Asn Leu Ile Asn Asn Thr Gly Leu Phe Phe Ile Gly Ile Gly Lys
 2020 2025 2030
 Arg Asn Ser Ser Thr Asn Thr Gly Asn Ser Ser Asp Ile Val Asn Tyr
 2035 2040 2045
 Gly Gln Tyr Asp Ser Met Gln Trp Ser Phe Ala Arg Ser Val Pro Met
 2050 2055 2060
 Asp Tyr Gln Val Ala Ala Val Ser Lys Gly Cys Tyr Phe Tyr Gln Lys
 2065 2070 2075 2080
 Thr Ser Asp Val Phe Asn Ser Glu Gly Met Tyr Pro Ser Asp Gly Gln
 2085 2090 2095
 Gly Met Gln Phe Val Asn Cys Ser Thr Asp His Leu Thr Met Phe Ser
 2100 2105 2110
 Val Gly Ala Phe Asn Pro Thr Ile Asp Ala Asp Phe Ser Tyr Asn Tyr
 2115 2120 2125
 Asn Val Asn Glu Ile Glu Lys Asn Val Lys Val Met Ile Ala Ala Val
 2130 2135 2140
 Phe Met Leu Val Val Tyr Gly Cys Leu Thr Ile Asn Ala Ile Ile Cys
 2145 2150 2155 2160
 Gln Arg Lys Asp Ala Ser Arg Gly Arg Leu Arg Phe Leu Lys Asp Asn
 2165 2170 2175
 Glu Pro His Asp Gly Tyr Met Tyr Val Ile Ala Val Glu Thr Gly Tyr
 2180 2185 2190
 Arg Met Phe Ala Thr Thr Asp Ser Thr Ile Cys Phe Asn Leu Ser Gly
 2195 2200 2205
 Asn Glu Gly Asp Gln Ile Phe Arg Ser Phe Arg Ser Glu Glu Asp Gly
 2210 2215 2220
 Asn Trp Glu Phe Pro Phe Ser Trp Gly Thr Thr Asp Arg Phe Val Met
 2225 2230 2235 2240
 Thr Thr Ala Phe Pro Leu Gly Glu Leu Glu Tyr Met Arg Leu Trp Leu
 2245 2250 2255
 Asp Asp Ala Gly Leu Asp His Arg Glu Ser Trp Tyr Cys Asn Arg Ile
 2260 2265 2270
 Ile Val Lys Asp Leu Gln Thr Gln Asp Ile Tyr Tyr Phe Pro Phe Asn
 2275 2280 2285
 Asn Trp Leu Gly Thr Lys Asn Gly Asp Gly Glu Thr Glu Arg Leu Ala
 2290 2295 2300
 Arg Val Glu Tyr Lys Arg Arg Phe Leu Asp Glu Ser Met Ser Met His
 2305 2310 2315 2320
 Met Leu Ala Gln Thr Ile Ser Trp Phe Ala Met Phe Thr Gly Gly Gly
 2325 2330 2335
 Asn Arg Leu Arg Asp Arg Val Ser Arg Gln Asp Tyr Ser Val Ser Ile
 2340 2345 2350
 Ile Phe Ser Leu Val Val Val Ser Met Ile Ser Ile Thr Ile Leu Lys
 2355 2360 2365
 Ser Asp Asn Ser Ile Ile Ser Asp Ser Lys Ser Val Ser Glu Phe Thr

2370	2375	2380
Phe Thr Ile Lys Asp Ile Ala Phe Gly Val Gly Phe Gly Val Leu Ile 2385 2390 2395 2400		
Thr Phe Leu Asn Ser Leu His Ile Leu Leu Cys Thr Lys Cys Arg Ser 2405 2410 2415		
His Ser Glu His Tyr Tyr Tyr Lys Lys Arg Lys Arg Glu Asp Pro Glu 2420 2425 2430		
Phe Lys Asp Asn Ser Gly Ser Trp Pro Met Phe Met Ala Gly Met Ala 2435 2440 2445		
Arg Thr Ile Ile Val Phe Pro Val Leu Met Gly Leu Ile Tyr Ile Ser 2450 2455 2460		
Gly Ala Gly Met Ser Leu Met Asp Asp Leu Ala Asn Ser Phe Tyr Ile 2465 2470 2475 2480		
Arg Phe Leu Ile Ser Leu Ile Leu Trp Ala Val Val Phe Glu Pro Ile 2485 2490 2495		
Lys Gly Leu Ile Trp Ala Phe Leu Ile Leu Lys Thr Arg Lys Ser His 2500 2505 2510		
Lys Ile Ile Asn Lys Leu Glu Glu Ala Leu Leu Arg Ala Lys Pro Ala 2515 2520 2525		
Glu Thr Phe Leu Arg Asn Pro Tyr Gly Lys Ile Glu Lys Gly Leu Gly 2530 2535 2540		
Thr Glu Ile Ala Asp Val Thr Lys Leu Arg Asp Thr Glu Asn Arg Lys 2545 2550 2555 2560		
Met Arg Asp Glu Gln Leu Phe Ile Thr Ile Arg Asp Met Leu Cys Phe 2565 2570 2575		
Phe Ala Ser Leu Tyr Ile Met Val Met Leu Thr Tyr Tyr Cys Lys Asp 2580 2585 2590		
Arg His Gly Tyr Trp Tyr Gln Leu Glu Met Ser Thr Ile Leu Asn Ile 2595 2600 2605		
Asn Gln Lys Asn Tyr Gly Asp Asn Thr Phe Met Ser Ile Gln His Ala 2610 2615 2620		
Asp Asp Phe Trp Asp Trp Ala Arg Glu Ser Leu Ala Thr Ala Leu Leu 2625 2630 2635 2640		
Ala Ser Trp Tyr Asp Gly Asn Pro Ala Tyr Gly Met Arg Ala Tyr Met 2645 2650 2655		
Asn Asp Lys Val Ser Arg Ser Met Gly Ile Gly Thr Ile Arg Gln Val 2660 2665 2670		
Arg Thr Lys Lys Ser Ala Glu Cys Thr Met Phe Lys Gln Phe Gln Gly 2675 2680 2685		
Tyr Ile Asn Asp Cys Gly Glu Glu Leu Thr Ser Lys Asn Glu Glu Lys 2690 2695 2700		
Thr Leu Tyr Met Gln Ala Gly Trp Thr Glu Leu Glu Ser Glu Asn Gly 2705 2710 2715 2720		
Thr Asp Ala Ser Asp Glu Tyr Thr Tyr Lys Thr Ser Glu Glu Leu Ser		

2725	2730	2735
Thr Glu Thr Val Ser Gly Leu Leu Tyr Ser Tyr Ser Gly Gly Gly Tyr		
2740	2745	2750
Thr Ile Ser Met Ser Gly Thr Gln Ala Glu Ile Ile Thr Leu Phe Asn		
2755	2760	2765
Lys Leu Asp Ser Glu Arg Trp Ile Asp Asp His Thr Arg Ala Val Ile		
2770	2775	2780
Ile Glu Phe Ser Ala Tyr Asn Ala Gln Ile Asn Tyr Phe Ser Val Val		
2785	2790	2795
Gln Leu Leu Val Glu Ile Pro Lys Ser Gly Ile Tyr Leu Pro Asn Ser		
2805	2810	2815
Trp Val Glu Ser Val Arg Leu Ile Lys Ser Glu Gly Ser Asp Gly Thr		
2820	2825	2830
Val Val Lys Tyr Tyr Glu Met Leu Tyr Ile Phe Phe Ser Val Leu Ile		
2835	2840	2845
Phe Val Lys Glu Ile Val Phe Tyr Leu Tyr Gly Arg Tyr Lys Val Ile		
2850	2855	2860
Thr Thr Met Lys Pro Thr Arg Asn Pro Phe Lys Ile Val Tyr Gln Leu		
2865	2870	2875
Ala Leu Gly Asn Phe Ser Pro Trp Asn Phe Met Asp Leu Ile Val Gly		
2885	2890	2895
Ala Leu Ala Val Ala Ser Val Leu Ala Tyr Thr Ile Arg Gln Arg Thr		
2900	2905	2910
Thr Asn Arg Ala Met Glu Asp Phe Asn Ala Asn Asn Gly Asn Ser Tyr		
2915	2920	2925
Ile Asn Leu Thr Glu Gln Arg Asn Trp Glu Ile Val Phe Ser Tyr Cys		
2930	2935	2940
Leu Ala Gly Ala Val Phe Phe Thr Ser Cys Lys Met Ile Arg Ile Leu		
2945	2950	2955
Arg Phe Asn Arg Arg Ile Gly Val Leu Ala Ala Thr Leu Asp Asn Ala		
2965	2970	2975
Leu Gly Ala Ile Val Ser Phe Gly Ile Ala Phe Leu Phe Phe Ser Met		
2980	2985	2990
Thr Phe Asn Ser Val Leu Tyr Ala Val Leu Gly Asn Lys Met Gly Gly		
2995	3000	3005
Tyr Arg Ser Leu Met Ala Thr Phe Gln Thr Ala Leu Ala Gly Met Leu		
3010	3015	3020
Gly Lys Leu Asp Val Thr Ser Ile Gln Pro Ile Ser Gln Phe Ala Phe		
3025	3030	3035
Val Val Ile Met Leu Tyr Met Ile Ala Gly Ser Lys Leu Val Leu Gln		
3045	3050	3055
Leu Tyr Val Thr Ile Ile Met Phe Glu Phe Glu Glu Ile Arg Asn Asp		
3060	3065	3070
Ser Glu Lys Gln Thr Asn Asp Tyr Glu Ile Ile Asp His Ile Lys Tyr		

3075

3080

3085

Lys Thr Lys Arg Arg Leu Gly Leu Leu Glu Pro Lys Asp Phe Ala Pro
 3090 3095 3100

Val Ser Ile Ala Asp Thr Gln Lys Asp Phe Arg Leu Phe His Ser Ala
 3105 3110 3115 3120

Val Ala Lys Val Asn Leu Leu His His Arg Ala Thr Arg Met Leu Gln
 3125 3130 3135

Thr Gln Gly Gln Tyr Gln Asn Gln Thr Val Ile Asn Tyr Thr Leu Ser
 3140 3145 3150

Tyr Asp Pro Val Ser Ala Ile His Glu Thr Gly Pro Lys Arg Phe Gln
 3155 3160 3165

Lys Trp Arg Leu Asn Asp Val Glu Lys Asp
 3170 3175

<210> 5

<211> 8073

<212> DNA

<213> C. Elegans *pkd-2* gene

<400> 5

tcattcttct tttttgtcag caatcgaggt gattgttgga cgacgagcgg cagattcacg 60
 gttacggact tgggttgtga ggagggcctg gacaagtaaa atatttattg gaaatttaga 120
 tatttagcag taacagcaaa attatttgta tttgttgtt taatttacta aatagtaaaa 180
 attgtaagtt ttcattaatt cttattgcca gaataaaaaa ttttctaatt ttgttttgtc 240
 taatttgtct aaaactacga aagtttttct ctaaaaattt cactagataa atacaatttt 300
 tcatgtttca attactttcc aaaagaagta acactataat tgcattagtt acaattttca 360
 actcacactc aaatccatca aatttcctcc atcttgttgt tgaactcttt gtttttcgat 420
 tgtctggagt gttgcattga ctccctcaat ccgatccaca atgctgaaca ctgattcttg 480
 catttgatct acacggcggg tcaaactgaa atgatttacg taatgtttat gatcatttat 540
 gatagagctg atacagtaaa agttaccaat tttgtttct attcttcgga attgtgaaaa 600
 aatacaattt tctcatgggt ttcattattt gaaaattcca gtcttcacac gtataaactg 660
 gaacacgaaa aactatgggt tttattctag aatactaatt ttttaatcga taaataatat 720
 tatcgtcaaa aaagcataaa gttttttttg taagatatat gaaaatcgaa taacaaaagt 780
 taaacttaat caatttatga aaacattgaa ccagtcacaaa atctaattgt gataccgtga 840
 aaaaaaacg tttccctcca aaagtttacc tttttcaagt cttctgttaa caaattttca 900
 gaacgtttat atttgtatgg tgacggtgaa acattatttg atcaaaaactg ctgtgggaac 960
 tgacggttat tatataatta aggttattat ggtaacagtg aaacagtatt taaaaatagc 1020
 tgtttcggta ctcaaggggt atcccatgag gaaaataaaa gtattacttt ttcagttatg 1080
 aaaactgaga atgttttcac aaaatgttac ctgtggtctg tttgggaaaa aggaaatcta 1140
 cgatgagaaa tttgcagaac attttttgtc aaaattctct acatgttttt tttgtttgta 1200

ggcagcacag cgggaagttca ggtgggttatg aaagagtaaa tatttttttt ctgtgatata 1260
 aaaaatgttt gcctgtcttg acggctgagg gccagcacat ttgcctacgt ttcaggtaaa 1320
 catgattttt gtaatttttc agtggcatgt agggccgcag gtaggcaggg ctacaaattt 1380
 gaccatttaa agttggttac acaataaaaat attaatcttc taaaatataa tcatttgaaa 1440
 attgaaatgc gaacottcgg ttattatoga attgaatgaa aaacaaaaag aaaataatto 1500
 taaaaactag ctgaacatc acaatttttc gtaaaactca ctttgogtaa tccctgtgat 1560
 tctccatata attctttttc tgttcogtca ttctagccac ctcatcagca atatcttcag 1620
 ccactttctc cgggaacatgc tcagtcattg atgtcacatc gaatcgagtg aatgcttcat 1680
 tgatgtcttt ttcagcgtat cgggcacggc agagcatcag ttgttagtct tcatacgtgg 1740
 cactctcttc aggggcctcc gggcgttttc caggttttgt gagtctctga actttctgga 1800
 atgaggatct ttgtggtttt agccaagcgc ccgacgtatt tggggaactc ttagaatatg 1860
 gggcgctgat gaacootgaa gcacccgaca tattccaggt tccaacataa acccagaaaa 1920
 tgtccgacgc tagtttaggt acaccaagta acttacatcc ataaaccaat ccaaaatccc 1980
 ctctccatct ttctttctag ccagctctgc ttccacttca acgtaggcat cattgatgat 2040
 agccaagaac atgttcaata ggatgaacga gacgaagaag acgtaggcaa tgaagaaggc 2100
 gggtcogaag aatcgattgc aggattctag agccgagaag ttaaagtcac ccgagaatgag 2160
 accggagcagg gogaacgcag agttgtagag gttggagtag tccggcatct tagaaaaatg 2220
 tgaagcgcgc gacattttac gggttttgtg taggcaaaaac ccggaagatg tcggatgcaa 2280
 gaatgtaaca tgcgatttag gatacttggt atcactcagt cagataacca ccatttttgt 2340
 taaaagaaaa tttactgttt cattcaagtt atataagtaa ttggaagatc ccgctgcggt 2400
 gaagcgtatt taagattggt aaaatagctg tgttgatatt tgggtacgtc aaaataaagg 2460
 aaatgaatgt tgtaatggat cagacatctg gcgggctcgg tgtaggcaga accaggcaga 2520
 tgtcggatgc atgaatgtaa aacgcctccg acatctgcgc ggttcttggt tcaaggtaag 2580
 cttgataata tttaaaaatg aaaaaaaaaa accaggcaga tgtcgggtgc taaaataatt 2640
 gctgcgaatt tcccgtttct taaactttat gagatggaaa tgaatcaaaa tgtcattgta 2700
 cctaagaatg cattogaatg gtagtaaaaa taaatgtttt tcatataaaa ttggtgaaac 2760
 tgcgattttt ttctaatttt atatttttta aatttcacag caatataaaa cgttacagta 2820
 cccaactat tctaaactcc acgaataaaa caaagatctt aaagattaag ttacctgtgt 2880
 cccaaagcac aaatatccaa actgtgcgaa tgcaaaaaag aaaacagcga acatcaactgc 2940
 aaatcctcca atatcctttg cagatctggt caacgtagag gacaactgtg acatgggtctt 3000
 gttaactgag atgaacttga acactttcac ccaagcaaca aataccacac atgctttgat 3060
 gttcagataa gagttctcgg aagaagtga gtcacgaat ggtgcatttg tcaatccgtt 3120
 ctcaatgaca gagttgacac gatttactcc ggtttttgtg cgattcactg acagaattat 3180
 tgtggctact gaaaatccta gcgcacaac gtctaccaa ttccagaact gggtagata 3240

gtggagacgg tgacggcoga tagcaaaaag ctcttcgaaa atgaagtata gtatgaatcc 3300
 acagaagatt ccttcaaaaa tcatcattcg ggtgcctcca gatgtttgat aggtcagaag 3400
 atcgtaagtc ataagctttg gagttgtgat aacacggcca gatgcaggga gctcaaatag 3420
 gagtctgaaa tgggaaattt cgaaaaaaat ttaactcgct gcttcagctt tatcataaaa 3480
 ttggcgcact tatttgaaaa ctattatctg atcgacattg attggaatgc aaatatattat 3540
 aataaatttg ttgacgtaac taaagtttta aaatccagtt taaaaaaact atgtaaaatt 3600
 tcagtactct tgaaactaga caagatttat acttgtttcc atttcocatag acacccctac 3660
 agttggcggg gtgactgata tgcattggcc gacatttttc gggttactgt ggattccatag 3720
 ttttcgggtg ggcccatctg aaggcaaaag tagtgcgggg cgaaaactcg aaaaacgtcg 3780
 accatgcata ccagtcacaa gccactcgaa ttccgaaatt ttggaatgaa cgtttactct 3840
 tgttgaaata cttataatta cagttctaca aacattgtta aatttttagt aaaaaagaga 3900
 caccattcca ccaaacatga tagaactcac ttcaccacac aaaaacagatt aatattcgca 3960
 ttgtacagag caaagtcac aataattgca cgtgatctct tctcgatcca ggcattagcc 4020
 tttaaagtg ccaattgcaga ttgagcttca gttgagccag ctaactggaag gcgttgaaac 4080
 aatccaccac ctccatatga agcaatggg cccacggttt ccaggtttcc aagctctttt 4140
 gccgtggcgt agatgaatct gaatataata ttttatttta aaaaaggatt ggtgagactg 4200
 ttttttatag gaatttatg ttgacaataa ctatctaaga ctaacaatta aatgaaaatt 4260
 gcattgacaa cataatgttc taaaatttaa aaaaaggagc atgaaacatt accaatatta 4320
 gttagaatat ttcaattttc gaggtacttt tcacaaactt tacatttttt tcaacgtttt 4380
 ttaataagaa tactctttca ggtagttaat atataagcta aattttgcat ttgtgtattg 4440
 aagcttttgc aaaaacacat aaacagatat aactgataat ttcttggaac ataaaattgt 4500
 attttcatgc aaatttcgta acattcttcc aaatacagtt tcataatatt gttaaaaaga 4560
 aattgggggt ttctcaatag tcataaaaat tctaaatatt tttaaaataa aactaagtat 4620
 tttccgcaaa taagtcaagt ttgcaataa aatttactgt ttcacattat gatcaagttt 4680
 gcattcacaat aagaaataat agtaaaaatt ggttctccat gaaaaaaccc cataaatgcc 4740
 atgaaacaac gttagctccg cctttcacca atcgccgatt ggtcagcaga attcaaaagg 4800
 tactagaagc tgetgattca acgaccaaac ttggccgaat ttacaaaatt gacgtcactc 4860
 acgcattcaac acttccatca ccgaccatcg tcttactctc gagcttttcc tcataatttg 4920
 caaaacattc cttaatctcc cgctggaaac ttttcattac agtacacgag tcatttgtea 4980
 ctttcaacat tctgatccga ggttcccca gcaaacgatt ctcatagtag atcatattct 5040
 cgttatccgt cgaattggaa gtttccgtcc aatatatgcc aggtattagg acttgtagaa 5100
 gccactgaaa gtttgatttg aaggttttca tttaaaaatt gaggaaactt acatcccaaa 5160
 tattatccat tgaggtagaa gatccaaatg ctggagctcc ggaggcacgc gtgctcgcca 5220

caaacagggtc gctcattact ttggagtagt agtaagattg gatgctgttt tgggagaacg 5280
 caactgtaaa tttttgaatt tagaaaaaaa aaacccgtga agtgcgggt gctaactggg 5340
 cgtgctcgat atatcacagg attagccga ctacctgoga ggtgctgggc gaaacactag 5400
 atgaaaattt tacaagaaaa tgattttoga aaatacaaac atttggttaac attaatgtta 5460
 tttttaagtt gttaaagcaa aaataaatat tggaaatttg aaaatgtttt gttacaaaaa 5520
 ttctgctgtt ttgcttacta agtaaaccta acaaatataa ggtaaaaata gtatgtgaac 5580
 gtttcatgag gttattcaag tagtgtgga aaattaaaaa gtgtagaaaa attacgtcac 5640
 aactgtatta aaatacataa aaacatgtat tttaatacat ttgtgacgtc acaaatgtat 5700
 ttaaatatat ttgctatat tacttgatta aacctattaa caaagttgta ctogtaaaat 5760
 ttcagttgaa atgctcaaac tcaactaaag tgttgaggaa aaaaaataaa aatttaaaaa 5820
 aaaactgttc caacgttgta acaaatgttg taacgggttg tcttaaatag tattcggagg 5880
 attcagcctg caatggacag ttttcaaaag agaaaaattt aactaattgg aagccattta 5940
 atcaaaaatt atgaatttag agattacttt gaaaaatgta tgattctaaa cgtttctttt 6000
 gtgtttattt gcaaaattca aatataagtt ttctactttt caaaaacctt ttataaaaaa 6060
 ttagaaaatt aaacaatttt ccaaacacaa tttttcccg tactgcatta aagtaacaa 6120
 ataaattgga agattagtaa ctactttggc catagtgttt caaacaaaagt gtggttttta 6180
 tgatgctcac aataaatttt tgaatgccg gttgaaacat ttttgaaaaa ttataaaaca 6240
 cgaaatgaat attttgcagt tgatagttac aaatccctgc caaatctttt ttttcacaaa 6300
 cttgaatttt aagaaatttg ctaaaaaaaaa acttcgggtg tttcatacat gccatataat 6360
 ttgtaaaaat aaagtgaaaa tggattcgtc gtgtgtagtt tggccactca ctataaaaat 6420
 gctgattaag tatagttagt ggagaaactc ggaaattgtc ggccggcgtg gaaacctacc 6480
 ccaaaaccgg accgagtgcg tcgggtggtg ttaaaatcgg acgaccggac gcgattttgt 6540
 acagccctat ttgaaagtaa tgacgtcata ctactttca tacagaaatt aaatatctga 6600
 tacgttagat tttgggaaat aagcttgta caaaaaatga tgtggtttat ttctagaagt 6660
 ctactatgt agttggtaca caaaatatga aatttgtagc gtatgcttca tagcagttac 6720
 aaagtcgaga actatttgta cattaatttg accaacaac ttaccataaa ccagcacaat 6780
 caagaacaca gcgtatccac caacttccat aaacgaacgg gcagtcagct tgatctttcc 6840
 atccgatttc tegtgtccgg atgccagcaa ggcttgagaa aacgagattc cctctttttg 6900
 agccggattc ttctcttat cgtgctcata ctctcactg accatagaat ggtcaaacga 6960
 ggggccatgc tccgcagcgg cgaaccgctg cggtggtta gcccatcgt cgtccgcagc 7020
 gccgtagttc attgaagacg gctcgtgaa acagtagaaa atttgaatta aagttttgag 7080
 aaaagttgaa aatcgagagc tctgtagtgt aaaaactgga aaaatagagt cgaaaagagg 7140
 cgagctcgcg aaatccacgt cctcgtagct cttggagatg ccgcattgct aagagatttc 7200
 cgtagatact atgttttatg ggatttcacg tttttggttg gagacggttt tttgcataga 7260

aacggaaaaa tgatgcagga atagaaaacg aacatgattt gaaactgaaa accatcgact 7320
atacggcaca atcatactac atttatcggy ttattgaaac tgcaccccaa aagttttacaa 7330
tttaaattca cataccattt gaagataaca accaataaaa agacttcgaa aggcgggaaa 7440
tgttgtggtt tgttggtgta gtggttatca catctgtcta acacacagaa ggtcgggtgtt 7500
tcgagccgcg ccgagatcat aagttttttg tcaatcatta atattgatto atctgaatga 7550
aattgtaaaa ttctttgaag gtgtttctaa atattgaaat gttttttttt agatttcggt 7620
agtatataat ttttgaaaca tacatttttt ttctccaaat ttcaagtata ttctacgatt 7630
tttgaaaaat cccaaaaaatt gttaacatta aaattctgaa taaacgggtgg aaatttgtag 7740
ttctctcaaa ttctaaataa aaattgaacg aaatttgaga aatttcctgt ttcaaaaaact 7800
aaatgtotta ttttcagagt tcaacaatgc cttagagaaa gttggaaaaat gataatgttt 7850
gttagtatat tgagaatata atgcaagtga aacaattagt ttttttttcg ataacaatta 7920
tttaaaaaaa actactgttt caaatctttt attcaaccaa tcttgtaata aaagttcact 7980
tatcttctcc ctcttcctcc ataatgtatg cccctcttca aatggaaaaa atgatgtcgg 8040
ggggaggtcc tcccccctcc caagaccctc cat 8073

<210> 6
<211> 815
<212> PRT
<213> C. Elegans Pkd-2 protein

<400> 6
Met Glu Gly Arg Gly Glu Gly Glu Asp Leu Pro Pro Thr Ser Tyr Phe
1 5 10 15
Pro Phe Glu Glu Gly His Thr Leu Trp Met Lys Arg Glu Lys Ile Lys
20 25 30
His Leu Gln Arg Ile Leu Gln Phe His Ser Asp Glu Ser Ile Leu Met
35 40 45
Ile Asp Lys Lys Leu Met Ile Ser Gly Gly Leu Glu Pro Pro Thr Phe
50 55 60
Cys Val Leu Asp Arg Cys Asp Asn His Tyr Thr Thr Lys Pro Arg His
65 70 75 80
Leu Pro Pro Phe Glu Val Phe Leu Phe Val Val Ile Phe Lys Cys Glu
85 90 95
Pro Ser Ser Met Asn Tyr Gly Ala Ala Asp Glu Arg Trp Ala Asn Pro
100 105 110
Pro Gln Pro Val Ala Ala Ala Glu His Gly Pro Ser Phe Asp His Ser
115 120 125
Met Val Ser Glu Glu Tyr Glu His Asp Lys Lys Lys Asn Pro Ala Gln
130 135 140
Lys Glu Gly Ile Ser Phe Ser Gln Ala Leu Leu Ala Ser Gly His Glu
145 150 155 160

Lys	Ser	Asp	Gly	Lys	Ile	Lys	Leu	Thr	Ala	Arg	Ser	Phe	Met	Glu	Val
				155					170					175	
Gly	Gly	Tyr	Ala	Val	Phe	Leu	Ile	Val	Leu	Val	Tyr	Val	Ala	Phe	Ala
			180					185					190		
Gln	Asn	Ser	Ile	Gln	Ser	Tyr	Tyr	Tyr	Ser	Lys	Val	Met	Ser	Asp	Leu
		195					200					205			
Phe	Val	Ala	Ser	Thr	Gly	Ala	Ser	Gly	Ala	Pro	Ala	Phe	Gly	Ser	Cys
	210					215					220				
Thr	Ser	Met	Asp	Asn	Ile	Trp	Asp	Trp	Leu	Ser	Gln	Val	Leu	Ile	Pro
225					230					235					240
Gly	Ile	Tyr	Trp	Thr	Glu	Thr	Ser	Asn	Ser	Thr	Asp	Asn	Glu	Asn	Met
				245					250					255	
Ile	Tyr	Tyr	Glu	Asn	Arg	Leu	Leu	Gly	Glu	Pro	Arg	Ile	Arg	Met	Leu
			260					265					270		
Lys	Val	Thr	Asn	Asp	Ser	Cys	Thr	Val	Met	Lys	Ser	Phe	Gln	Arg	Glu
		275					280						285		
Ile	Lys	Glu	Cys	Phe	Ala	Asn	Tyr	Glu	Glu	Lys	Leu	Glu	Asp	Lys	Thr
	290					295					300				
Met	Val	Gly	Asp	Gly	Ser	Val	Asp	Ala	Phe	Ile	Tyr	Ala	Thr	Ala	Lys
305					310					315					320
Glu	Leu	Glu	Asn	Leu	Lys	Thr	Val	Gly	Thr	Ile	Ala	Ser	Tyr	Gly	Gly
			325						330					335	
Gly	Gly	Phe	Val	Gln	Arg	Leu	Pro	Val	Ala	Gly	Ser	Thr	Glu	Ala	Gln
			340					345					350		
Ser	Ala	Ile	Ala	Thr	Leu	Lys	Ala	Asn	Arg	Trp	Ile	Asp	Arg	Gly	Ser
		355					360					365			
Arg	Ala	Ile	Ile	Val	Asp	Phe	Ala	Leu	Tyr	Asn	Ala	Asn	Ile	Asn	Leu
	370					375					380				
Phe	Cys	Val	Val	Lys	Leu	Leu	Phe	Glu	Leu	Pro	Ala	Ser	Gly	Gly	Val
385					390					395					400
Ile	Thr	Thr	Pro	Lys	Leu	Met	Thr	Tyr	Asp	Leu	Leu	Thr	Tyr	Gln	Thr
			405						410					415	
Ser	Gly	Gly	Thr	Arg	Met	Met	Ile	Phe	Glu	Gly	Ile	Phe	Cys	Gly	Phe
			420					425					430		
Ile	Leu	Tyr	Phe	Ile	Phe	Glu	Glu	Leu	Phe	Ala	Ile	Gly	Arg	His	Arg
		435					440					445			
Leu	His	Tyr	Leu	Thr	Gln	Phe	Trp	Asn	Leu	Val	Asp	Val	Val	Leu	Leu
	450					455					460				
Gly	Phe	Ser	Val	Ala	Thr	Ile	Ile	Leu	Ser	Val	Asn	Arg	Thr	Lys	Thr
465					470					475				480	
Gly	Val	Asn	Arg	Val	Asn	Ser	Val	Ile	Glu	Asn	Gly	Leu	Thr	Asn	Ala
				485					490					495	
Pro	Phe	Asp	Asp	Val	Thr	Ser	Ser	Glu	Asn	Ser	Tyr	Leu	Asn	Ile	Lys
			500					505					510		
Ala	Cys	Val	Val	Phe	Val	Ala	Trp	Val	Lys	Val	Phe	Lys	Phe	Ile	Ser

515					520					525					
Val	Asn	Lys	Thr	Met	Ser	Gln	Leu	Ser	Ser	Thr	Leu	Thr	Arg	Ser	Ala
530						535					540				
Lys	Asp	Ile	Gly	Gly	Phe	Ala	Val	Met	Phe	Ala	Val	Phe	Phe	Phe	Ala
545					550				555						560
Phe	Ala	Gln	Phe	Gly	Tyr	Leu	Cys	Phe	Gly	Thr	Gln	Ile	Ala	Asp	Tyr
				555					570					575	
Ser	Asn	Leu	Tyr	Asn	Ser	Ala	Phe	Ala	Leu	Leu	Arg	Leu	Ile	Leu	Gly
			580					585					590		
Asp	Phe	Asn	Phe	Ser	Ala	Leu	Glu	Ser	Cys	Asn	Arg	Phe	Phe	Gly	Pro
		595					600					605			
Ala	Phe	Phe	Ile	Ala	Tyr	Val	Phe	Phe	Val	Ser	Phe	Ile	Leu	Leu	Asn
	610					615					620				
Met	Phe	Leu	Ala	Ile	Ile	Asn	Asp	Ser	Tyr	Val	Glu	Val	Lys	Ala	Glu
625					630					635					640
Leu	Ala	Arg	Lys	Lys	Asp	Gly	Glu	Gly	Ile	Leu	Asp	Trp	Phe	Met	Asn
				645					650					655	
Lys	Val	Arg	Gly	Leu	Thr	Lys	Arg	Gly	Lys	Arg	Pro	Asp	Ala	Pro	Gly
			660					665					670		
Glu	Asp	Ala	Thr	Tyr	Glu	Asp	Tyr	Lys	Leu	Met	Leu	Tyr	Arg	Ala	Gly
		675					680					685			
Tyr	Ala	Glu	Lys	Asp	Ile	Asn	Glu	Ala	Phe	Thr	Arg	Phe	Asn	Val	Thr
	690					695					700				
Ser	Met	Thr	Glu	His	Val	Pro	Glu	Lys	Val	Ala	Glu	Asp	Ile	Ala	Asp
705					710					715					720
Glu	Val	Ala	Arg	Met	Thr	Glu	Gln	Lys	Arg	Asn	Tyr	Met	Glu	Asn	His
				725					730					735	
Arg	Asp	Tyr	Ala	Asn	Leu	Asn	Arg	Arg	Val	Asp	Gln	Met	Gln	Glu	Ser
			740					745					750		
Val	Phe	Ser	Ile	Val	Asp	Arg	Ile	Glu	Gly	Val	Asn	Ala	Thr	Leu	Gln
			755				760					765			
Thr	Ile	Glu	Lys	Gln	Arg	Val	Gln	Gln	Gln	Asp	Gly	Gly	Asn	Leu	Met
		770				775					780				
Asp	Leu	Ser	Ala	Leu	Leu	Thr	Asn	Gln	Val	Arg	Asn	Arg	Glu	Ser	Ala
785					790					795					800
Ala	Arg	Arg	Pro	Thr	Ile	Thr	Ser	Ile	Ala	Asp	Lys	Lys	Glu	Glu	
				805					810					815	

<210> 7

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Outside primer for PCR screening of lov-1 genomic (sy582) deletion

<400> 7

ctctatttgt ggttcgttgg cg

22

<210> 8
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Outside primer for PCR screening of
lov 1 genomic (sy582) deletion

<400> 8
ggggttttcc gttttcatgg gg

22

<210> 8
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Nested primer for PCR screening of
lov 1 genomic (sy582) deletion

<400> 8
ctaggaccca tgcaacagcg ag

22

<210> 10
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Nested primer for PCR screening of
lov-1 genomic (sy582) deletion

<400> 10
aacgctgatt ggttcaagtg tg

22

<210> 11
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Outside primer for PCR screening of
pkd-2 genomic (sy606) deletion

<400> 11
cccctcgttt gaccattcta tgg

23

<210> 12
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Outside primer for PCR screening of
pkd-2 genomic (sy606) deletion

<400> 12

acgtgatcct ctgtcgatcc ag

22

<210> 13
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<221> Description of Artificial Sequence: Nested primer for PCR screening of
pkd-2 genomic (sy606) deletion

<400> 13
agatcaagct gactgcooigt to

22

<210> 14
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<221> Description of Artificial Sequence: Nested primer for PCR screening of
pkd-2 genomic (sy606) deletion

<400> 14
gatccagcga ttagccttta acg

23

<210> 15
<211> 2870
<212> PRT
<213> C. Elegans Lov-1 sy582 deletion protein

<400> 15

Met Val Leu Arg Phe Ser Pro Pro Phe Arg Phe Ser Thr Thr Ser Phe
1 5 10 15

Phe Ser Cys Cys Leu Phe Cys Ser Glu Phe Ile Phe Val Phe Arg Arg
20 25 30

Ile Phe Thr Lys Leu Leu Gln Asp Asn Leu Pro Ala His Trp Met Lys
35 40 45

Lys Ser Asn Phe Phe Val Leu Leu Leu Leu Ala Ile Ser Ala Ile Gln
50 55 60

Ile Asp Gly Leu His Tyr Gln Leu Leu Asp Gly Ile Ala Thr Phe Arg
65 70 75 80

Leu Asp Asn Asp Asp Thr Thr Ile Gly Gly Val Pro Arg Asn Ser Gln
85 90 95

Gly Val Val Lys Ile Lys Leu Ser Cys Gly Leu Asn Arg Leu Ser Val
100 105 110

Glu Asn Lys Val Thr Glu Val Ser Ser Leu Glu Leu Ile His Asn Cys
115 120 125

Ile Gln Thr Glu Thr Arg Leu Val Gly Leu Phe Leu Asn Ser Thr Trp
130 135 140

Ile Thr Leu Asn Glu Val Asn Asp Asp Asp Glu Ile Ser Ile Ala Val
145 150 155 160

Glu Ala Lys Tyr Glu Val Cys Tyr Asp Asp Gly Ile Asp Arg Cys Asp

165										170					175				
Gly	Ser	Leu	Trp	Trp	Leu	Gln	Val	Gly	Gly	Asn	Glu	Met	Ala	Leu	Leu				
			180					185					190						
Gly	Tyr	Arg	Glu	Lys	Cys	Glu	Ser	Gly	Glu	Ile	Asn	Glu	Glu	Tyr	Ala				
		195					200					205							
Arg	Arg	Met	Cys	Lys	Arg	Pro	Tyr	Arg	Ser	Glu	Lys	Ser	Thr	Ala	Ile				
		210				215					220								
Ser	Asp	Ser	Gln	Gly	Val	Tyr	Tyr	Asp	Gly	Gln	Val	Leu	Lys	Gly	Val				
	225				230					235					240				
Arg	Ala	Lys	Gln	Phe	Ser	Met	Arg	Thr	Ser	Gly	Ser	Pro	Thr	Leu	Arg				
				245					250					255					
Arg	Met	Lys	Arg	Asp	Ala	Gly	Asp	Asn	Thr	Cys	Asp	Tyr	Thr	Ile	Glu				
			260					265					270						
Ser	Thr	Ser	Thr	Ser	Thr	Thr	Thr	Pro	Thr	Thr	Thr	Thr	Val	Thr	Ser				
		275					280						285						
Thr	Val	Thr	Ser	Thr	Thr	Thr	Val	Pro	Thr	Ser	Thr	Ser	Thr	Val	Thr				
	290					295					300								
Thr	Ala	Met	Ser	Thr	Ser	Thr	Ser	Thr	Pro	Ser	Thr	Ser	Thr	Thr	Ile				
	305				310					315					320				
Glu	Ser	Thr	Ser	Thr	Thr	Phe	Thr	Ser	Thr	Ala	Ser	Thr	Ser	Thr	Ser				
				325					330					335					
Ser	Thr	Ser	Thr	Thr	Gln	Gln	Ser	Ser	Ser	Thr	Ile	Thr	Ser	Ser	Pro				
			340					345					350						
Ser	Ser	Thr	Thr	Leu	Ser	Thr	Ser	Ile	Pro	Thr	Thr	Thr	Thr	Pro	Glu				
		355					360					365							
Ile	Thr	Ser	Thr	Leu	Ser	Ser	Leu	Pro	Asp	Asn	Ala	Ile	Cys	Ser	Tyr				
	370					375					380								
Leu	Asp	Glu	Thr	Thr	Thr	Ser	Thr	Thr	Phe	Thr	Thr	Thr	Met	Leu	Thr				
	385				390					395				400					
Ser	Thr	Thr	Thr	Glu	Glu	Pro	Ser	Thr	Ser	Thr	Thr	Thr	Thr	Glu	Val				
				405					410					415					
Thr	Ser	Thr	Ser	Ser	Thr	Val	Thr	Thr	Thr	Glu	Pro	Thr	Thr	Thr	Leu				
			420				425						430						
Thr	Thr	Ser	Thr	Ala	Ser	Thr	Ser	Thr	Thr	Glu	Pro	Ser	Thr	Ser	Thr				
		435					440					445							
Val	Thr	Thr	Ser	Pro	Ser	Thr	Ser	Pro	Val	Thr	Ser	Thr	Val	Thr	Ser				
	450					455					460								
Ser	Ser	Ser	Ser	Ser	Thr	Thr	Val	Thr	Thr	Pro	Thr	Ser	Thr	Glu	Ser				
	465				470					475				480					
Thr	Ser	Thr	Ser	Pro	Ser	Ser	Thr	Val	Thr	Thr	Ser	Thr	Thr	Ala	Pro				
				485				490						495					
Ser	Thr	Ser	Thr	Thr	Gly	Pro	Ser	Ser	Ser	Ser	Ser	Thr	Pro	Ser	Ser				
			500					505					510						
Thr	Ala	Ser	Ser	Ser	Val	Ser	Ser	Thr	Ala	Ser	Ser	Thr	Gln	Ser	Ser				

515					520					525					
Thr	Ser	Thr	Gln	Gln	Ser	Ser	Thr	Thr	Thr	Lys	Ser	Glu	Thr	Thr	Thr
530						535					540				
Ser	Ser	Asp	Gly	Thr	Asn	Pro	Asp	Phe	Tyr	Phe	Val	Glu	Lys	Ala	Thr
545					550					555					560
Thr	Thr	Phe	Tyr	Asp	Ser	Thr	Ser	Val	Asn	Leu	Thr	Leu	Asn	Ser	Gly
				565					570					575	
Leu	Gly	Ile	Ile	Gly	Tyr	Gln	Thr	Ser	Ile	Glu	Cys	Thr	Ser	Pro	Thr
			590					595					590		
Ser	Ser	Asn	Tyr	Val	Ser	Thr	Thr	Lys	Asp	Gly	Ala	Cys	Phe	Thr	Lys
		595					600					605			
Ser	Val	Ser	Met	Pro	Arg	Leu	Gly	Gly	Thr	Tyr	Pro	Ala	Ser	Thr	Phe
	610					615					620				
Val	Gly	Pro	Gly	Asn	Tyr	Thr	Phe	Arg	Ala	Thr	Met	Thr	Thr	Asp	Asp
625				630						635					640
Lys	Lys	Val	Tyr	Tyr	Thr	Tyr	Ala	Asn	Val	Tyr	Ile	Gln	Glu	Tyr	Ser
			645						650					655	
Ser	Thr	Thr	Ile	Glu	Ser	Glu	Ser	Ser	Thr	Ser	Ala	Val	Ala	Ser	Ser
			660					665					670		
Thr	Ser	Ser	Thr	Pro	Ser	Thr	Pro	Ser	Ser	Thr	Leu	Ser	Thr	Ser	Thr
		675					680					685			
Val	Thr	Glu	Pro	Ser	Ser	Thr	Arg	Ser	Ser	Asp	Ser	Thr	Thr	Thr	Ser
	690					695					700				
Ala	Gly	Ser	Thr	Thr	Thr	Leu	Gln	Glu	Ser	Thr	Thr	Thr	Ser	Glu	Glu
705					710					715					720
Ser	Thr	Thr	Asp	Ser	Ser	Thr	Thr	Thr	Ile	Ser	Asp	Thr	Ser	Thr	Ser
				725					730					735	
Thr	Ser	Ser	Pro	Ser	Ser	Thr	Thr	Ala	Asp	Ser	Thr	Ser	Thr	Leu	Ser
			740					745						750	
Val	Asp	Gln	Phe	Asp	Phe	Ile	Leu	Asp	Ser	Gly	Leu	Ser	Trp	Asn	Glu
	755					760					765				
Thr	Arg	His	Asn	Glu	Asp	Ser	Ile	Asn	Ile	Val	Pro	Leu	Pro	Thr	Asn
	770					775					780				
Ala	Ile	Thr	Pro	Thr	Glu	Arg	Ser	Gln	Thr	Phe	Glu	Cys	Arg	Asn	Val
785					790					795					800
Ser	Thr	Glu	Pro	Phe	Leu	Ile	Ile	Lys	Glu	Ser	Thr	Cys	Leu	Asn	Tyr
				805					810					815	
Ser	Asn	Thr	Val	Leu	Asn	Ala	Thr	Tyr	Ser	Ser	Asn	Ile	Pro	Ile	Gln
			820					825					830		
Pro	Ile	Glu	Thr	Phe	Leu	Val	Gly	Ile	Gly	Thr	Tyr	Glu	Phe	Arg	Ile
		835					840					845			
Asn	Met	Thr	Asp	Leu	Thr	Thr	Met	Gln	Val	Val	Ser	His	Ile	Phe	Thr
	850					855					860				
Leu	Asn	Val	Val	Ala	Asp	Ser	Thr	Ser	Thr	Ser	Glu	Val	Thr	Ser	Thr

865		870		875		880
Thr Ser Thr Gly Ser Ser Ser Glu Ser Ser Ala Ile Ser Thr Thr Ser						
		885		890		895
Gly Ile Glu Ser Thr Ser Thr Leu Glu Ala Ser Thr Thr Asp Ala Ser						
		900		905		910
Gln Asp Ser Ser Thr Ser Thr Ser Asp Ser Gly Thr Thr Ser Asp Ser						
		915		920		925
Thr Thr Ile Asp Ser Ser Asn Ser Thr Pro Ser Thr Ser Asp Ser Ser						
		930		935		940
Gly Leu Ser Gln Thr Pro Ser Asp Ser Ser Ser Ala Ser Asp Ser Met						
		945		950		955
Arg Thr Thr Thr Val Asp Pro Asp Ala Ser Thr Glu Thr Pro Tyr Asp						
		965		970		975
Phe Val Leu Glu Asn Leu Thr Trp Asn Glu Thr Val Tyr Tyr Ser Glu						
		980		985		990
Asn Pro Phe Tyr Ile Thr Pro Ile Pro Asn Lys Glu Pro Gly Ala Leu						
		995		1000		1005
Thr Thr Ala Met Thr Cys Gln Cys Arg Asn Asp Ser Ser Gln Pro Phe						
		1010		1015		1020
Val Leu Leu Lys Glu Ser Asn Cys Leu Thr Glu Phe Gly Lys Asn Gly						
		1025		1030		1035
Ala Tyr Ser Ala Ser Val Ser Phe Asn Pro Met Thr Ser Phe Val Pro						
		1045		1050		1055
Ala Thr Gly Thr Tyr Glu Phe Leu Ile Asn Val Thr Asn Arg Ala Ser						
		1060		1065		1070
Gly Glu Ser Ala Ser His Ile Phe Thr Met Asn Val Val Leu Pro Thr						
		1075		1080		1085
Thr Thr Thr Glu Thr Pro Pro Thr Thr Val Ser Ser Ser Asp Asp Ala						
		1090		1095		1100
Gly Gly Lys Thr Gly Gly Thr Gly Ala Thr Gly Gly Thr Gly Gly Thr						
		1105		1110		1115
Gly Ser Gly Gly Ser Ala Thr Thr Leu Ser Thr Gly Asp Ala Val Arg						
		1125		1130		1135
Ser Thr Thr Ser Gly Ser Gly Ser Gly Gln Ser Ser Thr Gly Ser Gly						
		1140		1145		1150
Ala Gly Gly Ser Gly Thr Thr Ala Ser Gly Ser Gly Ser Gly Gly Ser						
		1155		1160		1165
Ser Gly Thr Gly Ser Asp Gly Val Asn Ser Gly Lys Thr Thr Ala Leu						
		1170		1175		1180
Asn Gly Asp Gly Thr Gly Ser Gly Thr Ala Thr Thr Pro Gly Ser His						
		1185		1190		1195
Leu Gly Asp Gly Gly Ser Thr Ser Gly Ser Gly Ser Asp Ser Asn Gly						
		1205		1210		1215
Ser Ser Gly Val Ser Thr Lys Ser Ser Ser Gly Ser Asp Thr Ser Gly						

1220	1225	1230
Ser Ser Asp Ser Ser Gly Ala Asn Gly Ala Phe Ser Ala Thr Ala Gln 1235 1240 1245		
Pro Ser Thr Arg Thr Thr Lys Thr Arg Ser Ser Leu Ala Thr Val Ser 1250 1255 1260		
Pro Ile Ser Ala Ala Glu Gln Ala Ile Ile Asp Ala Gln Lys Ala Asp 1265 1270 1275 1280		
Val Met Asn Gln Leu Ala Gly Ile Met Asp Gly Ser Ala Ser Asn Asn 1285 1290 1295		
Ser Leu Asn Thr Ser Ser Ser Leu Leu Asn Gln Ile Ser Ser Leu Pro 1300 1305 1310		
Ala Ala Asp Leu Val Glu Val Ala Gln Ser Leu Leu Ser Asn Thr Leu 1315 1320 1325		
Lys Ile Pro Gly Val Gly Asn Met Ser Ser Val Asp Val Leu Lys Thr 1330 1335 1340		
Leu Gln Asp Asn Ile Ala Thr Thr Asn Ser Glu Leu Ala Asp Glu Met 1345 1350 1355 1360		
Ala Lys Val Ile Thr Lys Leu Ala Asn Val Asn Met Thr Ser Ala Gln 1365 1370 1375		
Ser Leu Asn Ser Val Leu Ser Ser Leu Asp Leu Ala Leu Lys Gly Ser 1380 1385 1390		
Thr Val Tyr Thr Leu Gly Val Ser Ser Thr Lys Ser Lys Asp Gly Thr 1395 1400 1405		
Tyr Ala Val Ile Phe Gly Tyr Val Ile Ala Ser Gly Tyr Thr Leu Val 1410 1415 1420		
Ser Pro Arg Cys Thr Leu Ser Ile Tyr Gly Ser Thr Ile Tyr Leu Thr 1425 1430 1435 1440		
Gly Asp Thr Arg Ala Ser Tyr Lys Gln Leu Asp Gly Asp Thr Val Thr 1445 1450 1455		
Ala Asp Thr Met Leu Ala Ala Ala Ile Gly Ile Gln Gly Met Phe Ala 1460 1465 1470		
Thr Asn Gly Arg Thr Val Gln Val Glu Gln Asp Lys Ile Asp Asp Lys 1475 1480 1485		
Arg Ser Leu Val Ser Gly Asn Ile Met Ala Thr Met Ser Gly Val Gly 1490 1495 1500		
Asp Val Gln Ser Gly Glu Tyr Ser Tyr Asn Asp Met Tyr Val Thr Ala 1505 1510 1515 1520		
Trp Asn Val Thr Tyr Asp Asn Ser Thr Val Gly Ser Thr Ser Gln Lys 1525 1530 1535		
Asn Thr Ser Phe Ser Phe Asn Ile Pro Val Ser Glu Val Gln Tyr Ile 1540 1545 1550		
Leu Leu Ile Glu Ser Gly Thr Met Ile Lys Leu His Ser Thr Gln Asn 1555 1560 1565		
Ile Val Ser Arg Gly Leu Val Val Thr Ala Ser Tyr Gly Gly Val Thr 1570 1575 1580		

Tyr Thr Ile Thr Cys Thr Asn Gly Thr Gly Lys Phe Val Glu Val Asp
 1535 1590 1595 1600
 Thr Asp Asn Ala Ile Phe Ser Tyr Asn Ala Asp Ser Phe Thr Val Val
 1605 1610 1615
 Ala Ser Asp Gly Ser Ser Ala Ser Thr Val Lys Lys Leu Ile Gln Met
 1620 1625 1630
 Pro Ile Val Ile Glu Asn Val Asn Leu Ala Leu Phe Asn Gln Thr Thr
 1635 1640 1645
 Ser Pro Leu Val Phe Ser Asn Ala Gly Ser Tyr Ser Met Arg Met Val
 1650 1655 1660
 Leu Ser Pro Gln Asp Ile Gly Ile Pro Ala Val Ser Ala Leu Ser Gln
 1665 1670 1675 1680
 Thr Val Ser Ile Ser Thr Leu Ser Pro Thr Ala Ser Tyr Thr Lys Asp
 1685 1690 1695
 Asp Leu Gln Ser Leu Ile Lys Glu Gln Thr Leu Val Thr Val Ser Gly
 1700 1705 1710
 Thr Thr Ser Asn Ser Leu Leu Ser Ile Ala Gly Ser Leu Thr Ser Ala
 1715 1720 1725
 Leu Lys Ile Ala Leu Asp Asn Pro Leu Ser Ser Asp Leu Ala Ala Asn
 1730 1735 1740
 Leu Lys Tyr Ala Thr Asp Asn Tyr Asp Ser Leu Tyr Asn Val Leu Pro
 1745 1750 1755 1760
 Ser Asp Pro Asp Asn Ile Val Tyr Val Glu Glu Met Thr Ser Glu Glu
 1765 1770 1775
 Trp Ala Ala Tyr Val Thr Lys Met Phe Gln Lys Asn Ile Ala Lys Asn
 1780 1785 1790
 Leu Ala Asn Gln Leu Ala Ser Thr Leu Asp Thr Leu Glu Asn Thr Leu
 1795 1800 1805
 Ala Ala Arg Ala Ile Ala Thr Gly Asn Leu Pro Tyr Asp Tyr Ser Asn
 1810 1815 1820
 Ser Val Asp Gly Thr Gly Met Val Ile Val Ile Asp Asp Ala Ser Asn
 1825 1830 1835 1840
 Ile Val Gly Lys Thr Gln Asn Cys Glu Glu Trp Ala Phe Lys Leu Pro
 1845 1850 1855
 Ser Pro Ala Ser Thr Leu Asn Thr Ala Glu Ile Thr Asp Lys Thr Leu
 1860 1865 1870
 Ile Gln Val Gly Leu Val Cys Tyr Ala Thr Asn Pro Arg Thr Tyr Val
 1875 1880 1885
 Asp Asn Phe Asp Met Leu Ile Thr Ser Gly Ala Leu Glu Ala His Ile
 1890 1895 1900
 Lys Asp Glu Asn Gln Ile Ile Ile Pro Ile Thr Gly Thr Thr Ala Pro
 1905 1910 1915 1920
 Ile Tyr Val Asn Gly Arg Gly Ser Glu Asp Asp Ala Val Leu Thr Leu
 1925 1930 1935

Met Gln Gln Gly Asp Phe Ala Ser Tyr Gln Ile Leu Asp Leu His Ala
 1940 1945 1950
 Phe Arg Thr Thr Asn Trp Asn Asn Ser Leu Gln Val Glu Ile Ile Ala
 1955 1960 1965
 Ser Gln Asp Tyr Glu Ile Pro Asn Asn Asp Asp Thr Tyr Met Phe Ser
 1970 1975 1980
 Ser Phe Gln Ser Leu Pro Gly Pro Leu Glu Ser Asn His Glu Trp Ile
 1985 1990 1995 2000
 Phe Asp Leu Asn Thr Leu Asn Lys Thr Ser Asn Tyr Phe Val Thr Ala
 2005 2010 2015
 Gly Asn Leu Ile Asn Asn Thr Gly Leu Phe Phe Ile Gly Ile Gly Lys
 2020 2025 2030
 Arg Asn Ser Ser Thr Asn Thr Gly Asn Ser Ser Asp Ile Val Asn Tyr
 2035 2040 2045
 Gly Gln Tyr Asp Ser Met Gln Trp Ser Phe Ala Arg Ser Val Pro Met
 2050 2055 2060
 Asp Tyr Gln Val Ala Ala Val Ser Lys Gly Cys Tyr Phe Tyr Gln Lys
 2065 2070 2075 2080
 Thr Ser Asp Val Phe Asn Ser Gln Gly Met Tyr Pro Ser Asp Gly Gln
 2085 2090 2095
 Gly Met Gln Phe Val Asn Cys Ser Thr Asp His Leu Thr Met Phe Ser
 2100 2105 2110
 Val Gly Ala Phe Asn Pro Thr Ile Asp Ala Asp Phe Ser Tyr Asn Tyr
 2115 2120 2125
 Asn Val Asn Glu Ile Glu Lys Asn Val Lys Val Met Ile Ala Ala Val
 2130 2135 2140
 Phe Met Leu Val Val Tyr Gly Cys Leu Thr Ile Asn Ala Ile Ile Cys
 2145 2150 2155 2160
 Gln Arg Lys Asp Ala Ser Arg Gly Arg Leu Arg Phe Leu Lys Asp Asn
 2165 2170 2175
 Glu Pro His Asp Gly Tyr Met Tyr Val Ile Ala Val Glu Thr Gly Tyr
 2180 2185 2190
 Arg Met Phe Ala Thr Thr Asp Ser Thr Ile Cys Phe Asn Leu Ser Gly
 2195 2200 2205
 Asn Glu Gly Asp Gln Ile Phe Arg Ser Phe Arg Ser Glu Glu Asp Gly
 2210 2215 2220
 Asn Trp Glu Phe Pro Phe Ser Trp Gly Thr Thr Asp Arg Phe Val Met
 2225 2230 2235 2240
 Thr Thr Ala Phe Pro Leu Gly Glu Leu Glu Tyr Met Arg Leu Trp Leu
 2245 2250 2255
 Asp Asp Ala Gly Leu Asp His Arg Glu Ser Trp Tyr Cys Asn Arg Ile
 2260 2265 2270
 Ile Val Lys Asp Leu Gln Thr Gln Asp Ile Tyr Tyr Phe Pro Phe Asn
 2275 2280 2285
 Asn Trp Leu Gly Thr Lys Asn Gly Asp Gly Glu Thr Glu Arg Leu Ala

2290	2295	2300
Arg Val Glu Tyr Lys Arg Arg Phe Leu Asp Glu Ser Met Ser Met His 2305 2310 2315 2320		
Met Leu Ala Gln Thr Ile Ser Trp Phe Ala Met Phe Thr Gly Gly Gly 2325 2330 2335		
Asn Arg Leu Arg Asp Arg Val Ser Arg Gln Asp Tyr Ser Val Ser Ile 2340 2345 2350		
Ile Phe Ser Leu Val Val Val Ser Met Ile Ser Ile Thr Ile Leu Lys 2355 2360 2365		
Ser Asp Asn Ser Ile Ile Ser Asp Ser Lys Ser Val Ser Glu Phe Thr 2370 2375 2380		
Phe Thr Ile Lys Asp Ile Ala Phe Gly Val Gly Phe Gly Val Leu Ile 2385 2390 2395 2400		
Thr Phe Leu Asn Ser Leu His Ile Leu Leu Cys Thr Lys Cys Arg Ser 2405 2410 2415		
His Ser Glu His Tyr Tyr Tyr Lys Lys Arg Lys Arg Glu Asp Pro Glu 2420 2425 2430		
Phe Lys Asp Asn Ser Gly Ser Trp Pro Met Phe Met Ala Gly Met Ala 2435 2440 2445		
Arg Thr Ile Ile Val Phe Pro Val Leu Met Gly Leu Ile Tyr Ile Ser 2450 2455 2460		
Gly Ala Gly Met Ser Leu Met Asp Asp Leu Ala Asn Ser Phe Tyr Ile 2465 2470 2475 2480		
Arg Phe Leu Ile Ser Leu Ile Leu Trp Ala Val Val Phe Glu Pro Ile 2485 2490 2495		
Lys Gly Leu Ile Trp Ala Phe Leu Ile Leu Lys Thr Arg Lys Ser His 2500 2505 2510		
Lys Ile Ile Asn Lys Leu Glu Gly Ser Asp Gly Thr Val Val Lys Tyr 2515 2520 2525		
Tyr Glu Met Leu Tyr Ile Phe Phe Ser Val Leu Ile Phe Val Lys Glu 2530 2535 2540		
Ile Val Phe Tyr Leu Tyr Gly Arg Tyr Lys Val Ile Thr Thr Met Lys 2545 2550 2555 2560		
Pro Thr Arg Asn Pro Phe Lys Ile Val Tyr Gln Leu Ala Leu Gly Asn 2565 2570 2575		
Phe Ser Pro Trp Asn Phe Met Asp Leu Ile Val Gly Ala Leu Ala Val 2580 2585 2590		
Ala Ser Val Leu Ala Tyr Thr Ile Arg Gln Arg Thr Thr Asn Arg Ala 2595 2600 2605		
Met Glu Asp Phe Asn Ala Asn Asn Gly Asn Ser Tyr Ile Asn Leu Thr 2610 2615 2620		
Glu Gln Arg Asn Trp Glu Ile Val Phe Ser Tyr Cys Leu Ala Gly Ala 2625 2630 2635 2640		
Val Phe Phe Thr Ser Cys Lys Met Ile Arg Ile Leu Arg Phe Asn Arg		

2645	2650	2655
Arg Ile Gly Val Leu Ala Ala Thr Leu Asp Asn Ala Leu Gly Ala Ile		
2660	2665	2670
Val Ser Phe Gly Ile Ala Phe Leu Phe Phe Ser Met Thr Phe Asn Ser		
2675	2680	2685
Val Leu Tyr Ala Val Leu Gly Asn Lys Met Gly Gly Tyr Arg Ser Leu		
2690	2695	2700
Met Ala Thr Phe Gln Thr Ala Leu Ala Gly Met Leu Gly Lys Leu Asp		
2705	2710	2715
Val Thr Ser Ile Gln Pro Ile Ser Gln Phe Ala Phe Val Val Ile Met		
2725	2730	2735
Leu Tyr Met Ile Ala Gly Ser Lys Leu Val Leu Gln Leu Tyr Val Thr		
2740	2745	2750
Ile Ile Met Phe Glu Phe Glu Glu Ile Arg Asn Asp Ser Glu Lys Gln		
2755	2760	2765
Thr Asn Asp Tyr Glu Ile Ile Asp His Ile Lys Tyr Lys Thr Lys Arg		
2770	2775	2780
Arg Leu Gly Leu Leu Glu Pro Lys Asp Phe Ala Pro Val Ser Ile Ala		
2785	2790	2795
Asp Thr Gln Lys Asp Phe Arg Leu Phe His Ser Ala Val Ala Lys Val		
2805	2810	2815
Asn Leu Leu His His Arg Ala Thr Arg Met Leu Gln Thr Gln Gly Gln		
2820	2825	2830
Tyr Gln Asn Gln Thr Val Ile Asn Tyr Thr Leu Ser Tyr Asp Pro Val		
2835	2840	2845
Ser Ala Ile His Glu Thr Gly Pro Lys Arg Phe Gln Lys Trp Arg Leu		
2850	2855	2860
Asn Asp Val Glu Lys Asp		
2865	2870	

<210> 16
 <211> 200
 <212> PRT
 <213> C. Elegans Pkd-2 deletion mutant (sy606) protein

<400> 16
 Met Glu Gly Arg Gly Glu Gly Glu Asp Leu Pro Pro Thr Ser Tyr Phe
 1 5 10 15
 Pro Phe Glu Glu Gly His Thr Leu Trp Met Lys Arg Glu Lys Ile Lys
 20 25 30
 His Leu Gln Arg Ile Leu Gln Phe His Ser Asp Glu Ser Ile Leu Met
 35 40 45
 Ile Asp Lys Lys Leu Met Ile Ser Gly Gly Leu Glu Pro Pro Thr Phe
 50 55 60
 Cys Val Leu Asp Arg Cys Asp Asn His Tyr Thr Thr Lys Pro Arg His
 65 70 75 80

Leu	Pro	Pro	Phe	Glu	Val	Phe	Leu	Phe	Val	Val	Ile	Phe	Lys	Cys	Glu
				85					90					95	
Pro	Ser	Ser	Met	Asn	Tyr	Gly	Ala	Ala	Asp	Glu	Arg	Trp	Ala	Asn	Pro
			100					105					110		
Pro	Gln	Pro	Val	Ala	Ala	Ala	Glu	His	Gly	Pro	Ser	Phe	Asp	His	Ser
			115				120					125			
Met	Val	Ser	Glu	Glu	Tyr	Glu	His	Asp	Lys	Lys	Lys	Asn	Pro	Ala	Gln
	130					135						140			
Lys	Glu	Gly	Ile	Ser	Phe	Ser	Gln	Ala	Leu	Leu	Ala	Ser	Gly	His	Glu
145					150				155					160	
Lys	Ser	Asp	Gly	Lys	Ile	Lys	Leu	Thr	Ala	Arg	Ser	Phe	Met	Glu	Val
			165					170						175	
Gly	Gly	Tyr	Ala	Val	Phe	Leu	Ile	Val	Leu	Val	Tyr	Asp	Ser	Ser	Thr
			180					185					190		
Pro	Arg	Gln	Lys	Ser	Leu	Lys	Thr								
			195				200								